Serial No.: 09/429,331

<u>REMARKS</u>

In the Office Communication dated June 13, 2008, the U.S. Patent and Trademark Office (hereinafter "the Patent Office") contends that the application does not comply with the requirements of 37 CFR § 1.821 through 1.825 pertaining to patent applications containing nucleotide sequence and/or amino acid sequence disclosures. In particular, the Patent Office contends that the specification recites sequences in the specification without an assigned SEQ ID No.

In response, applicants respectfully submit that a Response to Sequence Listing Requirement was filed on February 20, 2001, followed by a Supplemental Response to Sequence Listing Requirement on February 27, 2001, in response to an Office Communication of December 5, 2000. Similar to the instant Office Communication, the Office Communication of December 5, 2000 also alleged that the instant application did not comply with the requirements of 37 CFR § 1.821 through 1.825. The responses of February 20, 2001 and February 27, 2001 included amendments to the specification to add SEQ ID NOs. as well as sequence listings, both on paper and in computer readable format. Applicants submit herewith copies of the previously filed responses, as downloaded from the Patent Office Patent Application Information Retrieval (PAIR) system.

As such, applicants respectfully submit that the previously filed responses of February 20, 2001 and February 27, 2001 are believed to address the alleged deficiencies in the specification set forth in the instant Office Communication.

Accordingly, the instant application is believed to be in compliance with 37 CFR § 1.821 through 1.825.

Serial No.: 09/429,331

CONCLUSION

In light of the above remarks, it is respectfully submitted that the present application is now in proper condition for allowance, and an early notice to such effect is earnestly solicited.

If any small matter should remain outstanding after the Patent Examiner has had an opportunity to review the above remarks, the Patent Examiner is respectfully requested to telephone the undersigned patent attorney in order to resolve these matters and avoid the issuance of another Official Action.

DEPOSIT ACCOUNT

The Commissioner is hereby authorized to charge any additional fees associated with the filing of this correspondence to Deposit Account No. <u>23-1665</u>.

Respectfully submitted,

WIGGIN and DANA LLP

Date: 21 July 2008

By:

Todd E. Garabedian, Ph.D. Registration No. 39,197 Attorney for Applicants

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TADMITTED IN FL ONLY FRACTICE SUPERVISED BY FRINCIPALS OF THE FIRM

OL COÚ H26T IVER P. COOPER JAY M. FINKELSTEIN TELEFAX CONTROL SHEET

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aroly

DATE SENT:

SUBJECT:

No. of pages (including this cover sheet):

FROM:

Attached are the following

Remarks:

1) 2/20/01 Response to Sequence Listing...

and postcord receipt;

2) 2/27/01 Suppl. Resp. to Sequence Listing...

and postcord receipt.

CONFIDENTIALITY NOTE

This confidential facsimile message is intended only for the individual entity named above, and may contain information that is privileged and exempt from disclosure under applicable law. If you, the reader of this message, are not the intended recipient, or the employee or egent responsible for delivering this message to the intended recipient, you are hereby notified that you should not copy this facsimile or distribute it to envone other than the intended recipient. In addition, if you have received this telecopy in error, please immediately notify us by telephone or telefax and return the original message to us at the address above via the United States Postal Service. Finally, if it would not inconvenience you, we would appreciate it if you would first refax this message to the intended recipient. Thank you.

If this transmission is not well received, please advise us at our telecopier no. 202-737-3528 or by e-mail at mail@browdyneimark.com, or call our voice telephone no. 202-628-5197.

OR

OR

IN THE UNITED STATES PATENT AND TRADEMARK OFFICE

In Re Application of: PAIGE of at

Application No.: 09/429,331

Filed: October 28, 1999

For: METHOD OF PREDICTING THE ABILITY OF ...

Art Unit: 1627

Examiner: T. Wessendorf

Washington, D.C.

Ally.'s Docket: PAIGE=1D

Date: February 20, 2001

THE COMMISSIONER OF PATENTS AND TRADEMARKS Washington, D.C. 20231

Transmitted herewith is a [] Amendment [XX] Response to "Sequence Listing" Requirement with Sequence Listing and Disk and revised pages 239, 244-251, 266-268

in the above-identified application.

- [] Small Entity Status: Applicant(e) ctaim small entity status. See 37 C.F.R. §1.27.
- No additional foe is required. ()
- The fee has been calculated as shown below:

	(Col. 1)		(Col. 2)	(Cal. 3)		
CLAIMS REMAINING AFTER AMENDMENT			HIGHEST NO. PREVIOUSLY PAID FOR	PRESENT EXTRA EQUALS		
TOTAL	•	MINUS	20	D		
INDEP.		MINUS	3	0		
FIRST PR	FIRST PRESENTATION OF MULTIPLE DEP. CLAIM					

_	_	SINDAL	ENTITY
		RATE	ADDITIONAL FEE
1	x	9	\$
]	×	40	\$
j	1	135	S
ADDITION	NAL F	FEE TOTAL	s

SMALL ENTITY

OTHER THAN SMALL ENTITY RATE ADDITIONAL FEE 18 \$ 80 270 \$ TOTAL

- If the entry in Col. 1 is less than the entry in Col. 2, write "0" in Col. 3.
- If the "Highest Number Proviously Pald for" IN THIS SPACE is less than 20, write "20" in this space.
- If the "Highest Number Previously Paid for" IN THIS SPACE is less than 3, write "3" in this space.

The "Highest Number Previously Paid For" (lotal or independent) is the highest number found from the equivalent box in Col. 1 of a prior amendment of the number of claims originally filed.

Conditional Pathton for Extension of Time

If any extension of time for a response is required, applicant requests that this be considered a poblion therefor.

[XX] It is hereby politioned for an extension of time in accordance with 97 CFR 1.136(a). The appropriate fee required by 37 CFR 1.17 is calculated as shown below:

	Small Entity	Other Than Small Entity
	Responso Filed Within	Response Filed Within
	() First • \$ 55.00	[] First - \$ 110.00
	[] Second - \$ 195,00	(XX) Second - \$ 390,00
	[] Third - \$ 445.00	[] Third - \$ 890,00
	[] Fourth - \$ 695.00	[Fourth - \$ 1390.00
	Month After Time Period Set	Month After Time Period Set
1	[] Less (ees (\$) already paid formonth(s) exter Please charge my Deposit Account No. 02-4035 in the amount of	
ox]	Credit Card Payment Form, PTO-2038, Is attached, authorizing	
1	A check in the amount of \$ is attached (check)	
α	The Commissioner is hereby authorized and requested to charg overpayment to Deposit Account No. 02-4035. This authorization	o any additional foes which may be required in connec n and request is not limited to payment of all fees assu

[XX ction with this application or credit any overpayment to Deposit Account No. 02-4035. This authorization and request is not limited to payment of all fees associated with this communication, including studies and all palent processing fees under 37 CFR §1.17 throughout the prosecution of the case. This blanket authorization does not include patent studies patent studies patent studies and ell palent processing fees under 37 CFR §1.17 throughout the prosecution of the case. This blanket authorization does not include patent issue fees under 37 CFR §1.18.

BROWDY AND NEIMARK

Altomeys for Applicant(s)

er P. Coope Registration No. 28,005

Facsimile:

X Į

(202) 737-3528 (202) 828-6107

(NOVA) APPLICANT(S): PAIGE of al APPLICATION NO: 09/429,331	DOCKET NO.: PAIGE = 1 D
THE PATENT AND TRADEMARK OFFICE STAMP HEREON ACKNOWLEDGES RECEIPT OF THE FOLLOWING PAPERS: FEES \$ 390.00 PTO FORM 2038 (CH. #	CI FICO 26 Ph Chips
☐ MISSING PARTS RESPONSE WITH DECL ☐ AMENDMENT ☐ PRELIMINARY ☐ SUPPLEMENTAL ☐ REPLY TO OFFICE ACTION	ASSIGNMENT INFORMATION DISCLOSURE STATEMENT FORM 1449 & PATENTS/PUB: PRIORITY DOCUMENT(S) NO
RESTRICTION/ELECTION REPLY SEQUENCE LISTING WITH DISK RCE / CPA TRANSMITTAL (circle one) NOTICE OF APPEAL APPEAL BRIEF (TRIPLICATE) REPLY BRIEF (TRIPLICATE) OTHER TANAMA DAGES 239, ZYY	DECLARATION UNDER § LETTER TO DRAFTSMAN SHEETS OF DRAWINGS ISSUE FEE TRANSMITTAL FORM MAINTENANCE FEE LETTER 251, 266-268, 270 and 272
BEN-2 of Noted to Compli	HAND-CARRY

From-BROWDY NEIMARK

Application No.: (

NOTICE TO COMPLY WITH REQUIREMENTS FOR PATENT APPLICATIONS CONTAINING NUCLEOTIDE SEQUENCE AND/OR AMINO ACID SEQUENCE DISCLOSURES

The nucleotide and/or amino acid sequence disclosure contained in this application does not comply with the requirements for such a disclosure as set forth in 37 C.F.R. 1.821 - 1.825 for the following reason(s):

wing reason(s):
1. This application clearly fails to comply with the requirements of 37 C.F.R. 1.821-1.825. Applicant's attention is directed to these regulations, published at 1114 OG 29, May 15, 1990 and at 55 FR 18230, May 1, 1990.
2. This application does not contain, as a separate part of the disclosure on paper copy, a "S quence Listing" as required by 37 C.F.R. 1.821(c).
3. A copy of the "Sequence Listing" in computer readable form has not been submitted as required by 37 C.F.R. 1.821(e).
4. A copy of the "Sequence Listing" in computer readable form has been submitted. However, the content of the computer readable form does not comply with the requirements of 37 C.F.R. 1.822 and/or 1.823, as indicated on the attached copy of the marked-up "Raw Sequence Listing."
5. The computer readable form that has been filed with this application has been found to be damaged and/or unreadable as indicated on the attached CRF Diskette Problem Report. A Substitute computer readable form must be submitted as required by 37 C.F.R. 1.825(d).
6. The paper copy of the "Sequence Listing" is not the same as the computer readable form of the "Sequence Listing" as required by 37 C.F.R. 1.821(e).
7. Other:
Applicant Must Provide:
An initial or substitute computer readable form (CRF) copy of the "Sequence Listing".
An initial or substitute paper copy of the "Sequence Listing", as well as an amendment directing its entry into the specification.
A statement that the content of the paper and computer readable copies are the same and, where applicable, include no new matter, as required by 37 C.F.R. 1.821(e) or 1.821(f) or 1.821(g) or 1.825(b) or 1.825(d).
For questions regarding compliance to these requirements, please contact
For Rules Interpretation, call (703) 308-4216 For CRF Submission Help, call (703) 308-4212

For Patentin software help, call (703) 308-6856 PLEASE RETURN A COPY OF THIS NOTICE WITH YOUR RESPONSE

IN THE UNITED STATES PATENT AND TRADEMARK OFFICE

In re Application of:

PAIGE et al

Appln. No.: 09/429,331

Filed: October 28, 1999

For: METHOD OF PREDICTING THE ABILITY OF COMPOUNDS TO MODULATE THE BIOLOGICAL ACTIVITY OF RECEPTORS

ART UNIT: 1627

Examiner: T. Wessendorf

Washington, D.C.

February 20, 2001

Atty.Docket: PAIGE=1D

RESPONSE TO "SEQUENCE LISTING" REQUIREMENT

Honorable Commissioner of Patents Washington, D.C. 20231

sir:

In response to the Notice to Comply, mailed December 5, 2000, petition for a two-month extension of time and payment of late fees attached hereto, please amend the application as follows:

IN THE SPECIFICATION

Page 132, line 14, before "biotin" insert

--(SEQ ID NO:1)--
Page 136, line 26, after "...CTGCG" insert

--(SEQ ID NO:3)--;

line 27, after "...ACCTA" insert

--(SEQ ID NO:4)--.

In re Appln. No 09/429,331

Page 150, line 22, after "...GTCAG" insert -- (SEQ ID NO:5) --;

line 25, after "...GTCAG" insert

-- (SEQ ID NO: 6) --;

line 28, after "...GTCAG" insert

-- (SEQ ID NO:7)--;

line 31, after "...GTCAG" insert

-- (SEQ ID NO:8) --;

line 33, after "...TCGAG" insert

-- (SEQ ID NO:9) --.

Page 162, line 33, after "...CAGT-3'" insert

-- (SEQ ID NO:14) --;

line 36, after "...TAGA-3'" insert

-- (SEQ ID NO:15) --.

Page 173, line 26, after "...SLLSR" insert

-- (SEQ ID NO:187) --.

Page 183, line 6, after "SRLXXLL" insert

-- (SEQ ID NO:2) --.

Page 225, line 4, after "...KQAV" insert

-- (SEQ ID NO:10) --;

line 5, after "...GVSR" insert

-- (SEQ ID NO:11) --;

line 6, after "...MLSR" insert

-- (SEQ ID NO:12) --;

line 7, arter "...YASR" insert

May-15-2002 11:56

-- (SEQ ID NO:13) --.

Page 238, line 2, after "...GHSR" insert

-- (SEQ ID NO:59) --;

line 3, after "...WRSR" insert

-- (SEQ ID NO: 60) --;

line 4, after "...KDSR" insert

-- (SEQ ID NO: 61) --.

Attached are copies of pages 239, 244-251, 266-268, 270, and 272 in which sequence identifiers are marked in red. Entry of these revisions is respectfully requested.

Please enter the enclosed "Sequence Listing", pages 1-79.

REMARKS

- 1. Applicants hereby submit the following:
- [XX] a paper copy of a "Sequence Listing", complying with \$1.821(c), to be incorporated into the specification as directed above;
-] an amendment to the paper copy of the "Sequence
 Listing" submitted on , the amendment being in
 the form of substitute sheets;

- [XX] the Sequence Listing in computer readable form, complying with \$1.821(e) and \$1.824, including, if an amendment to the paper copy is submitted, all previously submitted data with the amendment incorporated therein;
- [] pursuant to \$1.821(e), reference is made to the computer readable form filed on , in USSN , which presents the identical Sequence information, the use of which is now requested, in lieu of submitting a new computer readable form; and/or
- [] a substitute computer readable form to replace one found to be damaged or unreadable.
- [XX] 2. The description has been amended to comply with \$1.821(d).
- 3. The undersigned attorney or agent hereby states as follows:
 - (a) this submission is not believed to include new matter [\$1.821(g)];

- (b) the contents of the paper copy (as amended, if applicable) and the computer readable form of the Sequence Listing, are believed to be the same [\$1.821(f) and \$1.825(b)];
- (c) if the paper copy has been amended, the amendment is believed to be supported by the specification and is not believed to include new matter [\$1.825(a)]; and
- (d) if the computer readable form submitted herewith is a substitute for a form found upon receipt by the PTO to be damaged or unreadable, that the substitute data is believed to be identical to that originally filed [\$1.825(d)].

Respectfully submitted,

BROWDY AND NEIMARK

Attorneys for Applicant(s)

By:

Gver P. Cooper

Registration No. 28,005

IPC:al
624 Ninth Street, N.W.
Washington, D.C. 20001
Telephone No.: (202) 628-5197
Facsimile No.: (202) 737-3528
F:\,\N\Nova\PaigeID\Pto\SequenceResponse.doc

239

Table 3: Phage/Peptide Classification # and isolation mathod E? + estradiol ER + estradiol #4 <u>class 1</u> SSNHQSSRLIELLSR 62 SRLKELLLPTDLSR 63 En + estradiol #15 SSKLYCLLDESYCSR 69 ER + estradiol #35 ER + estradiol 쓔수그 65 HGPLTLNLLRSSGG SRLEYWLKWEPGPSR 66 #12

class 2

SSCKWYEKCSGLWSR 67 #7 三民 ER + estradiol SSEYCFYWDSAHCSR 6 #33 <u>₹</u>3 #3Î #24 ER + estradiol 69 SSWVLLRDLPWGSR SSWVRLSDFPWGVSR 70 10

ER + estradiol Class 3 SSLTSRDFGSWYASR 71 #5

Class 4 15

ER SRTWESPLGTWEWSR 72 #13

Class 5

ER SAACATISHYLMGG 73 #48

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	present when present when peptide was jeptide was jeptide was ramoxifen Tamoxifen
244	Immobilized on Plastic No in the presence of the pep pep presence of the pep pep pep pep pep pep pep pep pep p
	Peptide Sequences Imme Peptide Sequence AN Peptide Sequence AN Peptide Sequence AN Peptide Sequence AN SERVICETWDDEYCSR 76 SERVICETWDDEYCSR 76 SERVICETWYDWYDWTFSR 80 SERVICETURGESR 80 SERVICET
	Table 7: New Ero Peptide name 1PT 2PT 3PT 4PT 5PT 10PT 11PT 12PT 13PT 13PT 14PT 15PT 16PT 18PT 18

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Estradiol Estradiol Estradiol Estradiol **Bstradio**l Estradiol gstradiol Estradiol Estradiol Raloxifen Raloxifen Tamoxifen Buffer Buffer Buffer Tamoxifen Tamoxifen Tamoxifen Buffer Buffer Buffer Estradio1 Buffer Buffer Estradiol Estradiol Estradiol present when peptide was identified H Table 8: New ER α -ERE Peptide Sequence Information מססט ਠ 8 ed in the W. presence of ಶ ℧ ಶ ರ receptor ਠ ช В form SRICFGDWCWLGGVDVLSR 109 SSINMVVDTPWCGKWVCSR 110 SSDWCFGWGGWCASEAVSR II7 SRNWDWAALELLPYPHPSR (K SRVFGVSGGEVVLINGSSR 10% SRLEELLLMDFWRSR (25 SKLPPSVFSMCGSEVCLSR 106 SSKLWQLLSSPIDSR 126 空主 SSKLYCLLDESYCSR [2] SRLTCLIASNGWDSEQCSR 103 SSLTSRDFGSWYASR 104 HSHNHHSPWLFRLLGG 100 SRFEIWKPEPGCVSSLENWE peptide Sequence 300 SSTGILWKLLTAESR SSHGILWRLLSEGSR SRSDSILWRMLSESR SRLVALLKSPWSVSR SSLTSRDFGSWYASR SSRPDAAFFGAKLSR SRSPILTHLLSLGSR SRSYHGEWGVWTLSR SSRIADLFWRLEPSR SRWNDTSWWLEELSR SSRPSESFWEKQLSR SSRPTAEWFRENLSR HSHPLPPLLSRLLTGG нзнрннзнггукгмдд SRTLQLDWGTLYSR PGKRVCSR 140 11.12 10B 김원원당 T1-10 T2-10 T3-11 14-10 **6B** 93 **SB** 1B 7.8 2R 3B E1-7 T3-1 4 B E1-4 Pepti K E1-1 E1-3 de name

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Estradiol Estradiol Estradiol Estradiol Estradiol Estradiol

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SRSLLMDMLMSDDYVTVSR 128
SSRLLACELMYEDADVCSR. 129
HSHSPLLMALLAPPGG 130
RIEYYLRIGTYESR 131
SSCLREILLYGACSR 132
SSRTAEDYCFFADDYWCSR 133
SSLRCYLSSSKVDQWACSR 135
SSLRCYLSSSKVDQWACSR 135
SSLRCYLSSKVDQWACSR 135
SSYKPHSLLEWHLLGGTSR 135

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Buffer Buffer Buffer Buffer Busser Busser Buffer Buffer Buffer Busser Buffer Buffer Buffer Buffer Buffer peptide was present when identified Table 9: New ERB-ERE Peptide Sequence Information Poplide Sequence Seq. of receptor [or SSEACYGRWMLCEQLGVSR 145 3 HSWLOPWRLSSIDLGG 157 133 Ē SSQVWPGPWRLVESR 146 SRGGGECLOPWCLSR 144 SSTSWLHYLMGTSR 143 SSGPWLMHYLGGGSR 142 SSGLPPNFERMLKSR 147 SSWPNPTFWERQLSR 138 HSYSSHPLLLSYLWGG SSKLYCLLDESYCSR SSWPGAEWFKEQLSR SSSLGRWRLSELESR SRECVGGWCLAELSR SSSOPWRWGLSIESR SSIPPRSWWLSQLSR SSSMMREFFERELSR SRLHCLLDSSYCSSR SRLHCLLDSSYCSSR SYSKEWFEERLNSR 23B-B 213-0 20Љ-В 17B-B 18B-B 193-6 16B-ß 14B-B 120-9 9B-B 7B-B 8B-B g-go 5B-β 4B-β 3B-6 2B-β 1B-B

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S

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Tamoxifen Tamoxifen Tamoxifen Tamoxifen Tamoxifen Tamoxifen Tamoxifen Fatradiol Tamoxifen Tamoxifen Tamoxifen Tamoxifen Tamoxifen Tamoxifen Tamoxifen Tamoxifen Buffer Tamoxifen Tamoxifen Tamoxifen Tamoxifen Tamoxifen SKIDLYSQMINEFFQINI.SR 175 SSRLVPHSFWLDGLMHGSR 173 SSISTYHMGEWFYAMLSSR 17# SRPGCLRGY.WCLADTPIPSR 172 SALNGVFCHDSSDLWYCSR 171 SSGPFYVGGMLWPADCLSR 169 SSDLINIDCLGVWCLSR 170 SRMFQVCGDEVCLRSR 169 Sregwmgpwrladsr 165 SSYASREWWYRELSK 168 SRLPPS.VFSMCGSEVCLSR (6) SSQEEWLLPWRLASR 162 SSARPWWLQFEGSSR 161 SSRGLLWDLLTKDSR SSPGSREWFKDMLSR SRNECIGPWCLTISR SREWEDGFGGRWLSR SRPYCLGDVWCLDSR HSTDMGWLRPWRLLGG SSTITMEDFFYERLSR SSWINSREFFLSQLSR SSVFTIMDGKVALSR 231.−β 22T-B 20T-β 18T-B 21T-B 17Τ-β 15T-B 16T-B 14T-B 10T-B 12T-β 13T-β 9T-B 8T-B 7T-A 6Τ-β 4T-β ST-β 2T-β lΤ-β

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Estradiol
SRLHDLLLRDESPSR 178 SRLHDLLLRDESPSR 179 SRDWRSGFLYELLSR 179 SSDTRSRLYELLSSSYTSR 181 SRLEELLRVGVLTSR 181 SSTGHRLLESLLLNSNSR 483 SSPTGHRLLESLLLNSNSR 483 SSPTGHRLLESLLLNSNSR 483 SSRTPILWHILLQDGSAETV 187 SSIKDFPNLISLLSR 186 SSIKDFPNLISLLSR 187 SSIKDFPNLISLLSR 187 SSIKDFPNLISLLSR 187 SSIKCLLNSNAFCSR 189 SSHCLLDAGQCSR 191 SSRLLCLLDQEACSR 192 SSLKCLLNSNFCSR 193 SSLKCLLNSNFCSR 193 SSLKCLLNSNFCSR 193 SSLKCLLNSNFCSR 193 SSLKCLLOSSPQKQPFCSR 194 SSRKLLEFIYLLGGSR 195 SSRKCLSEDMLRSRSR 196 SSRCLEDMLRSRSR 196 SSRCCSSLLGEMLIQTKESR 197 SSRCSSLLGEMLIQTKESR 197 SSLQAGSWLMHIYLRGGDSR 198
2E-B 3E-B 4E-B 5E-B 6E-B 6E-B 7E-B 10E-B 11E-B 12E-B 13E-B 14E-B 15E-B 17E-B 17E-B 19E-B 20E-B 20E-B

250.

Estradiol SSRPSFTIPWWFDDPSRSR 208 SSLDLSQFPMTASFLRESR 210 SSTRLMCWLGSADTSHCSR 203 SSYDWQCPSWYCPAPPSSR 204 SSYEIPKWALQWLSR 209 SSRTLYCHLTSSNPEWCSR 202 SSQWQAPWWYIDASR 201 SSWDFRVPWWYNNSR 2.6 SSTTWRCPEWYCGSR 205 SRWWLDDHELLLYSSR SSRTLLEHYLLGGSR SRPEGSSWLLHYLSR

276-β 286-β 29E-β 30E-β

2SE-β 26E-β 34E-B 35E-B

33E-B

31E-β 32E-β

Table 10: Panel Peptides for Example 2

Alternative name parenthesized. Modulator used to isolate peptide in brackets. BIII, SSEACVGRWMLCEQLGVSR. (B3) [no modulator] (Set 12 10 11) α/β III, SSWDMHQFFWEGVSR (AB3) [4-OH lamoxifen] (SEB 10 NO: 213) alp 1v, SRLPPSVFSMCGSEVCLSR (ABA) [same] (SEG 10 40:214) (SEE 10 MG; 211) B 11, SSLDLSQFPMTASFLRESK (B2) [17]2-estradiol] (Seg 15 rues 220) B 1, SREWEDGFGGRWISR (B1) [4-Ohl tamoxifen] (See 10 not 219) (372 KH 01 685) a III, SRTWESPLGTWEWSR (A3) [no modulator] (LEGG 13 PA): 21 8) α II, SSLTSRDFGSWYASR (A2) [17β-estradiol] (See 10 pol: 217) α 1, SSEYCFY WDSAHCSR (A1) [17fp-estradiol] (Seig 13 two: 216) (SER 18 MG: 212) a/B 11, SAPRATISHYLMGG (AB2) [no modulator] α/β 1, SSNHQSSRLIELLSR (AB1) [17 β -estradiol] α/β V, SSPGSREWFKDMLSR (ABS) {same}

S

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see is no:
         SRAGLĻSDLLEGKSR
 Table 100
                               223
         SSRSLLRDLLMVDSR
 A
         SSNKLLYNLLKMESR
                               224
          SSNKLLLNLLSTPSR
SSKSLLLNLLSTPSR
     HSFPRESLLVRLLQGG
              SRLEMLLRSETDFSR227
5.
               RLEELLKWGSVTSR 228
               RLEQLLKEEFSYSR229
               R'LEQLLRSEPDFSR230
               RLEDLLRAPFTTSR231
              S
               RLESLLRFGQLDSR 232
10
              S
               RLLSLLVGDFNSR
              s
               SRLEELLLGTNRDSR23Y
             S.S
               SRLEELLLMDFWRSR 235
                RLKELLLPTDL
                                 S R 236
                                 S R 237
               SRLECLLEGRLNC
15
               S
               SKLYCLLDESYCSR 238
                RLSCLLMGFEDCSR239
               SKLIRLLTSDE E.LSR 240
               SRLMELLQEGQGWSR 241
20
               SRLIELLSR 2/2
               SRLWQLLASTDTSR 26
       SSNHQS
               SRLWQLLSSPIDSR 244
SKLWQLLSSPIDSR 245
              s
                SRLVĀLLKSPWSVSR
              S
             SNSMLWKLLAAPSR 246
             SSKTLWRLLEGERSR 247
 25
             AGPVLWGLLSESR 248
             RSPILTHLLSLGSR 249
               TGILWKLLTAESR 250
             SHGILWRLLSEGSR251
 30
                  KLVQLLTTTAE 251
                  ILHRLLQEGSP 253
    В
                   LRYLLDKDEK 254
                  LLQQLLLTE 253
                  L
        SRCla
  35
                  QLSELLRGGSG 256
QLVLLLHAHKC 257
         CBP
                   YLEGLLMHQAA 258
                    LASLLQSESS259
                    LKTLLKKSKV260
   40
                    LALLISSEAH 261
                   H
                   Q
                    LLHLLKSQTI
                                  262
         RIP140
                   LLQLLLGHKNE 263
                   L
                   A T O'T T E U B K G 57%
                   LLSRLLRQNQD 265
                    VIKQLLLSENC 266
   45
          SRCla = human steroid receptor coactivator la,
                                      (CREB) -binding
                mouse cAMP-responsive element
          CBP
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protien, RIP 140 = human RIP140 50

May-15-2002 12:02

Table	101	CEO IB NO:
5	<u>(</u>	Class I ER4 SSNHQSRLIELLSR 267 D2 GSEPKSRLLELLSAPVTDV 280 D30 HPTHSSRLWELLMEATPTM 281 D11 VESGSSRLMQLLMANDLLT 282
		Class II
10		D47 HVYQHPLLLSLLSSEHESG 269 C33 HVEMHPLLMGLLMESQWGA 269 D14 QEAHGPLLWNLLSRSDTDW 270
10		Class III F6 GHEPLTLLERLLMDDKQAV 271 D22 LPYEGSLLLKLLRAPVEEV 272 D48 SGWENSILYSLLSDRVSLD 273 D43 AHGESSLLAWLLSGEYSSA 274 D43 AHGESSLLAWLLSGEYSSA 275
15		D17 GVFCDSILCUBLIAGSDAPS 776 D41 HHNGHSILYGLLAGSDAPS 77 D26 LGERASLLDMLLRQENPAW 217 D40 SGWNESTLYRLLQADAFDV 218 D40 SGWNESTLYRLLQADAFDV 217
. 20		D15 PSGGSSVLBIBBILD F4 PVGEPGLLWRLLSAPVERE 284
		Misc. D10 Weehsqmllhlldtgeavw6 283
E	ERβsp.	#293 SSIKDFPNLISLLSR [87
25	GRIP-1	NR1 DSKGQTKLLQLLTTKSDQM 16 NR2 LKEKHKILHQLLQDSSSPV 17 NR3 KKKENALLRYLLDKDDTKD 18
30	SRC-1	NR1 YSQTSHKLVKLLTTTAEQQ 19 NR2 LTARHKILHRLLQEGSPSD 20 NR3 ESKDHQLLRYLLDKDEKDL 21

		· 1T-	Peptic	les)
	. •	Sequence/Motif Aligned SEQUENCE/MOTIF Aligned	507	Library
	3 - 202A: Gial GDI	Sequence/Motif Aligned SPAHLLTWSEFLDSHTK	17.7	E
	Table ZVALLE	SRAHLLTWSEFLDSHTK SRAHLLTWSEFLDSHTK	BUF	-
	I <u>D</u>	SSGELITWYEFLGDLNP 23	50-	727
	<u>ID</u> 99	SSGELLTWIELSHGRP 24	BUF	, <u>E</u> K
	103	COCETATIVE	GTP	<u>K</u>
_	107	WELLED ~	GTP	CWL R Y
5		ALMADE TWENTY SP	BUF	R
	361	AMILIDERIOE		$\overline{\mathbf{v}}$
	388,391	NLMTWYEYLADGERL 28	GTP	PHD12
	45	NLMIWIDIA 29	BUF	
	397,401,412	*#####################################	GTP.	<u>N</u>
- 0	15r2.301,334	MAACI'A FE TED	BUF	· <u>H</u>
70	380,381,140			
	16	SSRGEIWWEI DO SSRGEIWWEI AREAGE 33	BOL	
	360	SSADG11 W		
	101	- There are	GTP	
	047	LGRGTTDMPPWAWWS 37	GDP	
15	375,123,125,247	MVTERPWVWIT	BUF	•
<u>.</u> -	331,334	WCMEDWKWYT JP	GTP)
	37	TATALCINE VIEC	GTP	
	- ·	EEGMDWFMRVVE38	Gřr	•
	387	- Direction of the Control of the Co		
	386			

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Mar-15-2002 12:02

		con-specific Phage				
	Table 202B: Glal	GTP-Specific Phage	GR ID NO):		
	(T-Peptides)		3 9 .	GTP	M	
	370,377,378	SVLSSSENCEGMACY	404		· <u>PARO</u> <u>K</u>	
5	244 366,G12	FNEVCLGWQCY SSNARPCQGWHCYLL	SQSR			
	G33,G34	wdggvwmgpas 4	٠ .	GTP	<u>X</u>	
	353 408	MGDSVLPYGGVWLGPEGNS ERVCGVWLGPEGNS	Y R 45	GTP	Ī	
10	G22,G25 G11,G26-29	SSWDGGVWWGQYGSR SSNLDGCFTSGGVWSGCSR	. 46 47			
10	G9,G10	LGYDINGVWIG	48	GTP	<u>N</u>	
	382	ICDIIPWEESCSR	49	GTP GTP	<u>P</u> PARO	
	384 413	ACGPAICPWDFMPQL	50	GIE	FARO	
	T 1 2					

Note: clone 244, which was identified in a screen for peptide which bound GDP:G-alpha, is suspected to having increased the affinity of the G-alpha for GTP through a conformational change.

	m was anace Gigal	GDP-Specific Phage				
	(D-Peptides)		ES IDAS	•		
		SRGPQLTWQEFLTGAASSR	গ্র			
	G4	NVVTVMEFLGP	52		*	-
	314	Srefvtwkeflgs	53	BUF	K	
5	73	SOLTWREFLFG	54	GD3	R	
	343	SSHLMTWHEFISD	55	GD B	H	
	217	. SRDGFETWAEFLGASGS	56	BUE		
	. 93	SRLTWSEYLSEIDP	51	BUF	CML	•
	62	SRTVTWVDFLKET	<i>5</i> 8	GD5	D	
10	193	MSWYEFMTEESM	SBZ	GDP	CWI	
	324	AKHDLSWYEFLQLPI	286	GTP	V	
	400	SRLSWWEFLGASDCGTC	787	GDP X1		
	281	DLLSLKEFLAT	288	GTP	K	
	359,161	SSPNLLTLEEFLS	287	GDP	L	
15	176	KTYSLYEFLEL	290	GTP	N	
	380,381,140	MSNRYTIYEFLNLHS	291	GTP	Y	
	409,24¥2	LHWWEVLAEK	292-	GDP	CWL	
	320	SSPQPLLHWWENMTEPP	293	GD₽		
	²³⁰	SRAGESVHWWEVL	294	GDP	H	
20	213	RAGPSEHWWEYIATL	295	GDP	N	
	266	EMISWHQYLLSI	ENN 296	•	GDP	PARO
	237		A 297	BUF	M	
	126,128,133,242	VPWWVWLAEGD	298	\mathtt{GTP}	N	٠
	379	SREIYWWDWLTDT	299	GD₽	. D	
25		FGSNMLDLPTFLD%L	300		PAR	0
	117	SRITFWELMLEGG	301	Buf	L	
	92	SRTPYEWLGYWGA	302	GDP	L	
	179				*** 4	OL.
		YDMCTWLEFLDGGEC	Z 03			CW
	289	SPLCTWAEYLMEPSC	70E			
30		TOWCTWAEFLSSTDC	305			
	273	SSDGCTWQEFLAGHGPC	306	GDE	. N	
	272,282,6R2		_		_	
•	225	PPNNPPWMWWS	30			
	337,339	SSPTVHENLPPWLWWSP	30	_		
_	268	LIHVPPNAWYD	30			
3		. GEDABBMAMD E	3 t			4CW
	329 280	YSQVFGDAPVWAWYSSR	311			
	319	WTPSDWQWWRSK	317		-	RO
	115	SSHWSSDSIFPGFWYSG	31	3 BU	r PP	7/0
	113		_	ur GD	D 90	
	10 197	SRGGVDLDIGNSA	31	4		
Ž,	197 347	EGEDVRTRIAN	31	5 GT	ie K	
	ジ ザ /	·				

SEQUENCE LISTING

<110> PAIGE, Lisa A. MCDONNELL, Donald P. CHANG, Ching Yu NORRIS, John HAMILTON, Paul T. FOWLKES, Dana M. BARNETT, Tom CHRISTIANSEN, Dale J. BUEHRER, Benjamin

- <120> METHOD OF PREDICTING THE ABILITY OF COMPOUNDS TO MODULATE THE BIOLOGICAL ACTIVITY OF RECEPTORS .
- <130> PAIGE1D
- <140> 09/429,331
- <141> 1999~10-28
- <150> PCT/US99/06664
- <151> 1999-03-26
- <150> 60/082,756
- <151> 1998-04-23
- <150> 60/099,656
- <151> 1998-09-09
- <150> 60/115,345
- <151> 1999-01-08
- <160> 315
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Gly ser Gly Lys

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Mat-15-2002 12:03

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<400> 3 gatctaggtc acagtgacct gcg	23
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<210> 7 <211> 73 <212> DNA <212> Arrificial Seguence	

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<223> Description of Artificial: Selected sequence from
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gaetgtgcga attetettet tgggatatgc atcaattttt ttgggaaggt gtttctagac 60
togagogtgt cag
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                 5
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                  5
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        occurrence is C or T
  agtgtgtgcc tegagannkn nknnknnknn knnknnkctg nnknnkctgc tgnnknnknn 60
  knnknnknnk nnktctagac tgtgcagt
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Ala Glu Gln Gln

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actgcacagt ctaga
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 Asp Gln Met
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 Leu Lys Glu Lys His Lys Ile Leu His Gln Leu Leu Gln Asp Ser Ser
  Ser Pro Val
  <210> 18
  <211> 19
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   Thr Lys Asp
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 Pro
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 Lys Thr Tyr Ser Leu Tyr Glu Phe Leu Glu Leu
                                      10
                  · 5
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  <210> 31
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<210> 32

<211> 13 <212> PRT

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Ser Ser Ala Asp Gly Ile Phe Trp Trp Glu Tyr Ala Arg Glu Ala Gly
                                      10
Glu
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Leu Gly Arg Gly Thr Thr Asp Met Pro Pro Trp Ala Trp Trp Ser
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                                      10
                                                       15
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                                    10
                  5
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 Glu Glu Gly Met Asp Trp Phe Met Arg Val Val Glu
             5
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Phe Asn Glu Val Cys Leu Gly Trp Gln Cys Tyr
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Gln Ser Arg
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Ser Ser Trp Asp Gly Gly Val Trp Trp Gly Gln Tyr Gly Ser Arg
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<211> 19
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Cys Ser Arg
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<211> 11
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                   5
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 1
                                                           15
Ser Ser Arg
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. Ser
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  <400> 57
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<400> 64

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                                       10
                                                           15
 <210> 72
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  Ser Arg Trp His Gly Thr Leu Phe Trp Gln Asp Glu Gln Ser Arg
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<211> 15

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<210> 77
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<400> 77
Ser Ser Cys Lys Trp Tyr Glu Lys Cys Ser Gly Leu Trp Ser Arg
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<210> 78
<211> 15
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Arbitrary
      peptide.
<400> 78
Ser Ser Arg Met Gly His Val Trp Tyr Asp Trp Thr Phe Ser Arg
<210> 79
<211> 15
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: Arbitrary
<400> 79
Ser Ser Arg Leu Leu Gly Asp Phe Gly Gly Ser Val Val Ser Arg
<210> 80
<211> 15
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: Arbitrary
      peptide
<400> 80
Ser Ser Lys Tyr Val Phe Gly Phe Gln Val Ala Gly Gly Ser Arg
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 <210> 81
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<212> PRT

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<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Arbitrary
      peptide
<400> 81
Ser Ser Trp Ala Gly Ile Lys Phe Gly Lys Pro Pro His Ser Arg
                                      10
<210> 82
<211> 15
<212> PRT
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<400> 82
Ser Ser Ser Trp Ser Tyr Gly Lys Pro Thr Phe Leu Ser Ser Arg
                                      10
<210> 83
<211> 15
<212> PRT
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      peptide
<400> 83
Ser Arg Asp Thr Gly Asp Met Trp Trp Gly Arg Gly Gly Ser Arg
                                      10
<210> 84
<211> 15
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     peptide
<400> 84
Ser Ser Gly Arg Tyr Asp Pro Phe Val Leu Asn Ala Ala Ser Arg
                                     10
                   5
<210> 85
<211> 15
<212> PRT
<213> Artificial Sequence
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<400> 85
Ser Ser Ser Pro Trp Trp Ser Phe Asn Leu Arg Asp Met Ser Arg
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<210> 86
<211> 15
<212> PRT
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<223> Description of Artificial Sequence: Arbitrary
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<400> 86
Ser Ser Trp Pro Tyr Leu Pro Lys Arg Glu Glu Trp Ala Ser Arg
<210> 87
<211> 15
<212> PRT
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<223> Description of Artificial Sequence: Arbitrary
     peptide
<400> 87
ser Ser Gly Trp Ile Glu Gln Lys Leu Arg Gly Ser Phe Ser Arg
<210> 88
<211> 15
<212> PRT
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<223> Description of Artificial Sequence: Arbitrary
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<400> 88
Ser Ser Ser Ala Thr Ser Ile Lys Val Gln Tyr Gln Ile Ser Arg
<210> 89
<211> 15
<212> PRT
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<220>
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 <400> 89
 Ser Ser Tyr Leu Thr Leu Gly Lys Ser Met Met Ala Ile Ser Arg
                                      10
  <210> 90
  <211> 15
  <212> PRT
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  <220>
  <223> Description of Artificial Sequence: Arbitrary
       peptide
  <400> 90
  Ser Ser Trp His Ser Arg Trp Asp Leu Ala Leu Gly Phe Ser Arg
                    5
  <210> 91
  <211> 15
  <212> PRT
  <213> Artificial Sequence
  <223> Description of Artificial Sequence: Arbitrary
        peptide
   <400> 91
   Ser Ser Gly Tyr Trp Gly Gly Trp Asp Tyr Gly Ala Gly Ser Arg
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   <210> 92
   <211> 15
   <212> PRT
   <213> Artificial Sequence
   <223> Description of Artificial Sequence: Arbitrary
         peptide
   <400> 92
   Ser Arg Asp Asn Cys Gly Ala Gly Leu Trp Ala Gly Cys Ser Arg
                     S
   <210> 93
  · <211> 15
   <212> PRT
   <213> Artificial Sequence
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   <223> Description of Artificial Sequence: Arbitrary
         peptide
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<400> 93 ·
Ser Ser Ser Thr Pro Gly Trp Trp Glu Trp Asp Trp Ala Ser Arg
                                     10
<210> 94
<211> 19
<212> PRT
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      peptide
Ser Ser Tyr Trp Asp Gly Ser Trp Arg Arg Lys Glu Thr Cys Val Ser
                                     10
Cys Ser Arg
<210> 95
<211> 19
<212> PRT
<213> Artificial Sequence
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<223> Description of Artificial Sequence: Arbitrary
      peptide
Ser Ser Arg Thr Ala Glu Asp Tyr Cys Phe Phe Ala Asp Asp Tyr Trp
                                      10
Cys Ser Arg
<210> 96
 <211> 15
 <212> PRT
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       peptide
 <400> 96
 Ser Ser Arg Ala Leu Ala Leu Phe Pro Val Gly Met Glu Ser Arg
                  5
 <210> 97
 <211> 19
 <212> PRT
 <213> Artificial Sequence
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<223> Description of Artificial Sequence: Arbitrary
     peptide
<400> 97
Ser Ser Asp Cys Glu Ser Leu Thr Ser Tyr Pro His Leu Lys Ala Leu
                                      10
Cvs Ser Arg
<210> 98
<211> 15
<212> PRT
<213> Artificial Sequence
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<223> Description of Artificial Sequence: Arbitrary
     peptide
<400> 98
Ser Ser Thr Ala Thr Ala Leu Arg Asp Arg Leu Ala Tyr Ser Arg
                                      10
<210> 99
<211> 15
<212> PRT
 <213> Artificial Sequence
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      peptide
 <400> 99
 Ser Ser Gly Lys Thr Arg Glu His Tyr Arg Glu Gly Thr Ser Arg
                  5
                                      10
 <210> 100
 <211> 16
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 <400> 100
 His Ser His Ash His His Ser Pro Trp Leu Phe Arg Leu Leu Gly Gly
                                       10
 <210> 101
 <211> 16
 <212> PRT
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<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Arbitrary
      peptide
His Ser His Pro His His Ser His Leu Leu Tyr Lys Leu Met Gly Gly
                   5
  1
 <210> 102
 <211> 16
 <212> PRT
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 <220>
 <223> Description of Artificial Sequence: Arbitrary
       peptide
 <400> 102
 His Ser His Pro Leu Pro Pro Leu Leu Ser Arg Leu Leu Thr Gly Gly
                                       10
                   5 ·
 <210> 103
 <211> 19
 <212> PRT
<213> Artificial Sequence
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       peptide
 <400> 103
 Ser Arg Leu Thr Cys Leu Leu Gln Ser Asn Gly Trp Asp Ser Glu Gln
 Cys Ser Arg
  <210> 104
  <211> 15
  <212> PRT
  <213> Artificial Sequence
  <223> Description of Artificial Sequence: Arbitrary
        peptide
  <400> 104
  Ser Ser Leu Thr Ser Arg Asp Phe Gly Ser Trp Tyr Ala Ser Arg
                                        10
  <210> 105
<211> 14
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<212> PRT

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<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Arbitrary
      peptide
<400> 105
Ser Arg Thr Leu Gln Leu Asp Trp Gly Thr Leu Tyr Ser Arg
<210> 106
<211> 19
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: Arbitrary
      peptide
Ser Arg Leu Pro Pro Ser Val Phe Ser Met Cys Gly Ser Glu Val Cys
Leu Ser Arg
<210> 107
<211> 28
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: Arbitrary
      peptide
Ser Arg Phe Glu Ile Trp Lys Pro Glu Pro Gly Cys Val Ser Ser Leu
                                      10
Glu Asn Trp Glu Pro Gly Lys Arg Val Cys Ser Arg
                                  25
             20
<210> 108
<211> 19
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: Arbitrary

    peptide

Ser Arg Val Phe Gly Val Ser Gly Gly Glu Val Val Leu Ile Asn Gly
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Ser Ser Arg
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<210> 109

<211> 19

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:Arbitrary
 peptide ..

<400> 109

Ser Arg Leu Cys Phe Gly Asp Trp Cys Met Leu Gly Gly Val Asp Val 1 5 10 15

Leu Ser Arg

<210> 110

<211> 19

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:Arbitrary
 peptide

<400> 110

Ser Ser Leu Asn Met Val Val Asp Thr Pro Trp Cys Gly Lys Trp Val 1 5 10 15

Cys Ser Arg

<210> 111

<211> 15

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:Arbitrary
 peptide

<400> 111

Ser Ser Arg Pro Asp Ala Ala Phe Phe Gly Ala Lys Leu Ser Arg 1 5 10 15

<210> 112

<211> 15

<212> PRT

<213> Artificial Sequence

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<223> Description of Artificial Sequence: Arbitrary
     peptide
<400> 112
Ser Ser Arg Pro Ser Pro Ser Phe Trp Glu Lys Gln Leu Ser Arg
                                     10
<210> 113
<211> 15
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Arbitrary
<400> 113
Ser Ser Arg Pro Thr Ala Glu Trp Phe Arg Glu Asn Leu Ser Arg
<210> 114
<211> 15
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Arbitrary
      peptide
<400> 114
Ser Arg Trp Trp Asp Thr Ser Trp Trp Leu Glu Glu Leu Ser Arg
<210> 115
<211> 15
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: Arbitrary
      peptide
<400> 115
Ser Ser Arg Ile Ala Asp Leu Phe Trp Arg Leu Glu Pro Ser Arg
<210> 116
<211> 15
 <212> PRT
<213> Artificial Sequence
 <223> Description of Artificial Sequence: Arbitrary
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peptide
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<400> 116
Ser Arg Ser Tyr His Gly Glu Trp Gly Val Trp Thr Leu Ser Arg
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<210> 117

<211> 19

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Arbitrary peptide

<400> 117

ser Ser Asp Trp Cys Phe Gly Trp Gly Gly Trp Cys Ala Ser Glu Ala 10

Val Ser Arg

<210> 118

<211> 19

<212> PRT

<213> Artificial Sequence

<223> Description of Artificial Sequence: Arbitrary peptide

<400> 118

Ser Arg Asn Trp Asp Trp Ala Ala Leu Glu Leu Leu Pro Tyr Pro His 10

Pro Ser Arg

<210> 119

<211> 15

<212> PRT

<213> Artificial Sequence

<223> Description of Artificial Sequence: Arbitrary peptide

<400> 119

Ser Ser Leu Thr Ser Arg Asp Phe Gly Ser Trp Tyr Ala Ser Arg

<210> 120

<211> 15

<212> PRT

<211> 15 <212> PRT

<213> Artificial Sequence

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<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Arbitrary
     peptide
<400> 120
Ser Arg Ser Pro Ile Leu Thr His Leu Leu Ser Leu Gly Ser Arg
                 5
                                     10
<210> 121
<211> 15
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Arbitrary
    peptide
<400> 121
Ser Ser Thr Gly Ile Leu Trp Lys Leu Leu Thr Ala Glu Ser Arg
                  5
                                     10
 1
<210> 122
<211> 15
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Arbitrary
     peptide
Ser Ser His Gly Ile Leu Trp Arg Leu Leu Ser Glu Gly Ser Arg
                                     10
<210> 123
<211> 15
<212> PRT
<213> Artificial Sequence
<220>
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     peptide
<400> 123
Ser Arg Ser Asp Ser Ile Leu Trp Arg Met Leu Ser Glu Ser Arg
<210> 124
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<220>
<223> Description of Artificial Sequence: Arbitrary
      peptide
<400> 124
Ser Arg Leu Val Ala Leu Leu Lys Ser Pro Trp Ser Val Ser Arg
<210> 125
<211> 15
 <212> PRT
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 <223> Description of Artificial Sequence: Arbitrary
       peptide
 <400> 125
 Ser Arg Leu Glu Glu Leu Leu Met Asp Phe Trp Arg Ser Arg
                                       10
 <210> 126
 <211> 15
<212> PRT
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 <223> Description of Artificial Sequence: Arbitrary
       peptide
 <400> 126
 Ser Ser Lys Leu Trp Gln Leu Leu Ser Ser Pro Ile Asp Ser Arg
                                      10
 <210> 127
 <211> 15
 <212> PRT
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 <400> 127
 Ser Ser Lys Leu Tyr Cys Leu Leu Asp Glu Ser Tyr Cys Ser Arg
                                       10
                    5
   1
  <210> 128
  <211> 19
  <212> PRT
  <213> Artificial Sequence
  <223> Description of Artificial Sequence: Arbitrary
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peptide
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Val Ser Arg

<210> 129

<211> 19

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Arbitrary peptide

<400> 129

Ser Ser Arg Leu Leu Ala Cys Glu Leu Met Tyr Glu Asp Ala Asp Val 1 5 10 15

Cys Ser Arg

<210> 130

<211> 16

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:Arbitrary
 peptide

<400> 130

His Ser His Ser Pro Leu Leu Met Ala Leu Leu Ala Pro Pro Gly Gly
1 10 15

<210> 131

<211> 15

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:Arbitrary
 peptide

<400> 131

Ser Arg Leu Glu Tyr Tyr Leu Arg Leu Gly Thr Tyr Glu Ser Arg 1 5 10

<210> 132

<211> 15

<212> PRT

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<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Arbitrary
      peptide
·<400> 132
Ser Ser Cys Leu Arg Glu Ile Leu Leu Tyr Gly Ala Cys Ser Arg
                                      10
<210> 133
<211> 19
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Arbitrary
      peptide
<400> 133
Ser Ser Arg Thr Ala Glu Asp Tyr Cys Phe Phe Ala Asp Asp Tyr Trp
Cys Ser Arg
<210> 134
<211> 19
<212> PRT
<213> Artificial Sequence
 <223> Description of Artificial Sequence: Arbitrary
      peptide
 <400> 134
Ser Ser Leu Arg Cys Tyr Leu Ser Ser Ser Lys Val Asp Gln Trp Ala
                                                           15
                  5
 Cys Ser Arg
 <210> 135
 <211> 19
 <212> PRT
 <213> Artificial Sequence
 <220>
 <223> Description of Artificial Sequence: Arbitrary
     peptide
 <400> 135
 Ser Ser Tyr Lys Pro His Ser Leu Leu Glu Trp His Leu Leu Gly Gly
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```
Thr Ser Arg
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<210> 136
<211> 15
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Arbitrary
     peptide
<400> 136
Ser Arg Leu His Cys Leu Leu Asp Ser Ser Tyr Cys Ser Ser Arg
<210> 137
<211> 15
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: Arbitrary
      peptide
<400> 137
Ser Arg Leu His Cys Leu Leu Asp Ser Ser Tyr Cys Ser Ser Arg
                                      10
<210> 138
<211> 15
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Arbitrary
      peptide
<400> 138
Ser Ser Trp Pro Asn Pro Thr Phe Trp Glu Arg Gln Leu Ser Arg
                   5
                                      10
 1 .
<210> 139
<211> 14
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: Arbitrary
    · peptide
<400> 139
 Ser Tyr Ser Lys Glu Trp Phe Glu Glu Arg Leu Asn Ser Arg
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<210> 140
<211> 15
<212> PRT
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 <223> Description of Artificial Sequence: Arbitrary
      peptide
 <400> 140
 Ser Ser Ser Met Met Arg Glu Phe Phe Glu Arg Glu Leu Ser Arg
 <210> 141
 <211> 15
 <212> PRT
 <213> Artificial Sequence
 <220>
 <223> Description of Artificial Sequence: Arbitrary
       peptide
 <400> 141
 Ser Ser Gly Leu Pro Pro Asn Phe Glu Arg Met Leu Lys Ser Arg
                                       10
 <210> 142
 <211> 15
 <212> PRT
 <213> Artificial Sequence
 <220>
 <223> Description of Artificial Sequence: Arbitrary
      peptide
 <400> 142
  Ser Ser Gly Pro Trp Leu Met His Tyr Leu Gly Gly Gly Ser Arg
                                       10
  <210> 143
  <211> 15
  <212> PRT
  <213> Artificial Sequence
  <223> Description of Artificial Sequence: Arbitrary
       peptide
  <400> 143
  Ser Ser Thr Ser Trp Leu His His Tyr Leu Met Gly Thr Ser Arg
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<210> 144
<211> 15
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Arbitrary
     peptide
<400> 144
Ser Arg Gly Gly Glu Cys Leu Gly Pro Trp Cys Leu Ser Arg
                                     10
<210> 145
<21.1> 19
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<223> Description of Artificial Sequence: Arbitrary
      peptide
<400> 145
Ser Ser Glu Ala Cys Val Gly Arg Trp Met Leu Cys Glu Gln Leu Gly
                                      10
Val Ser Arg
<210> 146
<211> 15
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: Arbitrary
      peptide
<400> 146
Ser Ser Gln Val Trp Pro Gly Pro Trp Arg Leu Val Glu Ser Arg
                                      10
<210> 147
<211> 15
<212> PRT
<213> Artificial Sequence
 <223> Description of Artificial Sequence: Arbitrary
     peptide
 <400> 147
 Ser Ser Ser Leu Gly Pro Trp Arg Leu Ser Glu Leu Glu Ser Arg
                   5
                                      10
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<210> 148

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<211> 15
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Arbitrary
      peptide
 <400> 148
 Ser Ser Ser Gly Pro Trp Arg Trp Gly Leu Ser Ile Glu Ser Arg
                                                          15
                  5 .
 <210> 149
 <211> 15
 <212> PRT
 <213> Artificial Sequence
 <223> Description of Artificial Sequence: Arbitrary
      peptide
 <400> 149
 Ser Arg Glu Cys Val Gly Gly Trp Cys Leu Ala Glu Leu Ser Arg
 <210> 150
 <211> 15
<212> PRT
 <213> Artificial Sequence
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      peptide
 Ser Ser Ile Pro Pro Arg Ser Trp Trp Leu Ser Gln Leu Ser Arg
                                       10
                                                            15
                  . 5
 <210> 151
 <211> 15
 <212> PRT
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 <223> Description of Artificial Sequence: Arbitrary
       peptide
 <400> 151
 Ser Ser Trp Pro Gly Ala Glu Trp Phe Lys Glu Gln Leu Ser Arg
 <210> 152
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<211> 15
<212> PRT
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<223> Description of Artificial Sequence: Arbitrary
      peptide
<400> 152
Ser Ser Lys Leu Tyr Cys Leu Leu Asp Glu Ser Tyr Cys Ser Arg
                                      10
<210> 153
<211> 16
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Arbitrary
      peptide
<400> 153
His Ser Tyr Ser Ser His Pro Leu Leu Leu Ser Tyr Leu Trp Gly Gly
                                                            15
                  5
                                       10
<210> 154
<211> 16
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Arbitrary
      peptide
<400> 154
His Ser Trp Leu Gly Pro Trp Arg Leu Ser Ser Ile. Asp Leu Gly Gly
<210> 155
<211> 16
<212> PRT
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 <223> Description of Artificial Sequence: Arbitrary
      peptide
 <400> 155
His Ser Thr Asp Met Gly Trp Leu Arg Pro Trp Arg Leu Leu Gly Gly
                                       10
 <210> 156
<211> 15
<212> PRT
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<211> 15 <212> PRT

<213> Artificial Sequence

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<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Arbitrary
<400> 156
Ser Ser Val Phe Thr Ile Met Asp Gly Lys Val Ala Leu Ser Arg
                 5
                                      10
<210> 157
<211> 15
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Arbitrary
      peptide
Ser Arg Pro Tyr Cys Leu Gly Asp Val Trp Cys Leu Asp Ser Arg
                  5
<210> 158
<211> 15
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Arbitrary
      peptide
<400> 158
Ser Arg Glu Trp Glu Asp Gly Phe Gly Gly Arg Trp Leu Ser Arg
.1
                                      10
<210> 159
<211> 15
<212> PRT
<213> Artificial Sequence
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<223> Description of Artificial Sequence: Arbitrary
      peptide
<400> 159
Ser Ser Trp Asn Ser Arg Glu Phe Phe Leu Ser Gln Leu Ser Arg
                 5
 1
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<210> .160
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<220>
 <223> Description of Artificial Sequence: Arbitrary
      peptide
 <400> 160
 Ser Ser Thr Thr Met Phe Asp Phe Phe Tyr Glu Arg Leu Ser Arg
                                       10
 <210> 161
 <211> 15.
 <212> PRT
 <213> Artificial Sequence
 <220>
 <223> Description of Artificial Sequence: Arbitrary
      peptide
 <400> 161
 Ser Ser Ala Arg Pro Trp Trp Leu Gln Phe Glu Gly Ser Ser Arg
                  5
 <210> 162
 <211> 15
 <212> PRT
 <213> Artificial Sequence ·
 <223> Description of Artificial Sequence: Arbitrary
      peptide
 Ser Ser Gln Glu Glu Trp Leu Leu Pro Trp Arg Leu Ala Ser Arg
                   5
                                       10
 <210> 163
 <211> 19
 <212> PRT
 <213> Artificial Sequence
  <223> Description of Artificial Sequence: Arbitrary
       peptide
  <400> 163
 Ser Arg Leu Pro Pro Ser Val Phe Ser Met Cys Gly Ser Glu Val Cys
 Leu Ser Arg
  <210> 164
  <211> 19
  <212> PRT
  <213> Artificial Sequence
```

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<223> Description of Artificial Sequence: Arbitrary
      peptide
<400> 164
Ser Ser Gly Pro Phe Tyr Val Gly Gly Met Leu Trp Pro Ala Asp Cys
                  5
  1
                                      10
Leu Ser Arg
<210> 165
<211> 15
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Arbitrary
     peptide ·
<400> 165
Ser Arg Glu Gly Trp Met Gly Pro Trp Arg Leu Ala Asp Ser Arg
  1
                                      10
<210> 166
<211> 15
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: Arbitrary
     peptide
<400> 166
Ser Arg Asn Glu Cys Ile Gly Pro Trp Cys Leu Thr Ile Ser Arg
                                      10
<210> 167
<211> 15 .
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Arbitrary
     peptide
<400> 167
Ser Ser Pro Gly Ser Arg Glu Trp Phe Lys Asp Met Leu Ser Arg
                                      10
<210> 168
<211> 15
<212> PRT
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<210> 172 <211> 19

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<213> Artificial Sequence
<223> Description of Artificial Sequence: Arbitrary
      peptide
<400> 168
Ser Ser Val Ala Ser Arg Glu Trp Trp Val Arg Glu Leu Ser Arg
                                      10
<210> 169
<211> 16
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Arbitrary
      peptide
<400> 169
Ser Arg Met Phe Gln Val Cys Gly Asp Glu Val Cys Leu Arg Ser Arg
                                      10
<210> 170
<211> 16
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: Arbitrary
     peptide
<400> 1.70
Ser Ser Asp Leu His Arg Asp Cys Leu Gly Val Trp Cys Leu Ser Arg
<210> 171
<211> 19
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Arbitrary
      peptide
<400> 171
Ser Arg Leu Asn Gly Val Phe Cys His Asp Ser Ser Asp Leu Trp Val
                                     10
                                                          15
Cys Ser Arg
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<212> PRT
  <213> Artificial Sequence
  <223> Description of Artificial Sequence: Arbitrary
        peptide
  <400> 172
  Ser Arg Pro Gly Cys Leu Arg Gly Val Trp Cys Leu Ala Asp Thr Pro
                     5
                                        10
                                                             15
  Pro Ser Arg
  <210> 173
  <211> 19
  <212> PRT
  <213> Artificial Sequence
  <220>
  <223> Description of Artificial Sequence: Arbitrary
        peptide
  <400> 173
. Ser Ser Arg Leu Val Pro His Ser Phe Trp Leu Asp Gly Leu Met His
                                        10
  Gly Ser Arg
  <210> 174
  <211> 19
  <212> PRT
  <213> Artificial Sequence
. <220>
  <223> Description of Artificial Sequence: Arbitrary
        peptide :
  <400> 174
  Ser Ser Ile Ser Thr Tyr His Met Gly Glu Trp Phe Tyr Ala Met Leu
                    5
                                       10
                                                            15
  Ser Ser Arg
  <210> 175
  <211> 18
  <212> PRT
  <213> Artificial Sequence
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  <223> Description of Artificial Sequence: Arbitrary
       peptide
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<223> Description of Artificial Sequence: Arbitrary
     peptide
<400> 187
Ser Ser Ile Lys Asp Phe Pro Asn Leu Ile Ser Leu Leu Ser Arg
<210> 188
<211> 19 ·
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Arbitrary.
      peptide
<400> 188
Ser Ser Gly Ser Ser Ala Gly Arg Leu Met Mer Leu Leu Gln Asp Gly
                 . 5
Val Ser Arg
<210> 189
<211> 15
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Arbitrary
      peptide
<400> 189
Ser Arg Glu Gly Leu Leu Met Arg Leu Leu Ile Gly Asp Ser Arg
                                      10
<210> 190
<211> 15
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Arbitrary
      peptide
<400> 190
Ser Ser His Cys His Thr Arg Leu Cys Ser Leu Leu Thr Ser Arg
                                                           15
                   5
<210>.191
<211> 15
<212> PRT
 <213> Artificial Sequence
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<223> Description of Artificial Sequence: Arbitrary
     peptide
<400> 191
Ser Ser Arg Leu Leu Cys Leu Leu Asp Ala Gly Gln Cys Ser Arg
                  5
<210> 192
<211> 15
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: Arbitrary
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<400> 192
Ser Arg Asn Leu Leu Cys Leu Leu Asp Gln Glu Ala Cys Ser Arg
    . 5
                                     10
<210> 193
<211> 14
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Arbitrary
     peptide
<400> 193
Ser Ser Leu Lys Cys Leu Leu Asn Ser Asn Phe Cys Ser Arg
<210> 194
<211> 19
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: Arbitrary
     peptide
<400> 194
Ser Ser Leu Lys Cys Leu Leu Gln Ser Ser Pro Gln Lys Gln Pro Phe
Cys Ser Arg
<210> 195
<211> 15
<212> PRT
<213> Artificial Sequence
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<223> Description of Artificial Sequence: Arbitrary
      peptide
<400> 195
Ser Ser Arg Thr Leu Leu Glu His Tyr Leu Leu Gly Gly Ser Arg
<210> 196
<211> 15
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Arbitrary
      peptide
<400> 196
Ser Ser Ala Gly Leu Leu Glu Asp Met Leu Arg Ser Arg Ser Arg
                                      10
<210> 197
<211> 19
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Arbitrary
      peptide
<400> 197
Ser Ser Arg Cys Ser Ser Leu Leu Cys Glu Met Leu Ile Gln Thr Lys
                  5
Glu Ser Arg
<210> 198
<211> 19
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Arbitrary
      peptide
<400> 198
Ser Ser Leu Gln Ala Gly Ser Trp Leu Met His Tyr Leu Arg Gly Gly
                                      10
                                                          15
Asp Ser Arg
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<210> 199
 <211> 15
 <212> PRT
 <213> Artificial Sequence
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 <223> Description of Artificial Sequence: Arbitrary
      peptide
 <400> 199
Ser Arg Pro Glu Gly Ser Ser Trp Leu Leu His Tyr Leu Ser Arg
                                                          15
<210> 200
<211> 15
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: Arbitrary
     peptide
<400> 200
Ser Ser Arg Thr Leu Leu Glu His Tyr Leu Leu Gly Gly Ser Arg
                                      10 .
<210> 201
<211> 16
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Arbitrary
     peptide
<400> 201
Ser Arg Trp Trp Leu Asp Asp His Glu Leu Leu Leu Tyr Ser Ser Arg
                5
                                      10
<210> 202
<211> 19
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: Arbitrary
     peptide
<400> 202
Ser Ser Arg Thr Leu Tyr Cys His Leu Thr Ser Ser Asn Pro Glu Trp
                  5
                                    10
Cys Ser Arg
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<210> 203
 <211> 19
 <212> PRT
 <213> Artificial Sequence
 <220>
 <223> Description of Artificial Sequence: Arbitrary
<400> 203
Ser Ser Thr Arg Leu Met Cys Trp Leu Gly Ser Ala Asp Thr Ser His
Cys Ser Arg
<210> 204
<211> 19
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: Arbitrary
      peptide
<400> 204
Ser Ser Tyr Asp Trp Gln Cys Pro Ser Trp Tyr Cys Pro Ala Pro Pro
Ser Ser Arg
<210> 205
<211> 15
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Arbitrary
    peptide
<400> 205
Ser Ser Thr Thr Trp Arg Cys Pro Glu Trp Tyr Cys Gly Ser Arg
                                      10
<210> 206
<211> 15
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Arbitrary
     peptide
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```
<400> 206
Ser Ser Trp Asp Phe Arg Val Pro Trp Trp Tyr Asn Asn Ser Arg
  ı
                   5
                                       10
<210> 207
<211> 15
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Arbitrary
<400> 207
Ser Ser Gln Trp Gln Ala Pro Trp Trp Tyr Ile Asp 'Ala Ser Arg
                                      10
<210> 208
<211> 19
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Arbitrary
      peptide
Ser Ser Arg Pro Ser Phe Thr Ile Pro Trp Trp Phe Asp Asp Pro Ser
                  5
                                      10
Arg Ser Arg
<210> 209
<211> 15
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Arbitrary
     peptide
Ser Ser Tyr Glu Ile Pro Lys Trp Ala Leu Gln Trp Leu Ser Arg
                 5
<210> 210
<211> 19
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Arbitrary
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Ser Ser Leu Asp Leu Ser Gln Phe Pro Met Thr Ala Ser Phe Leu Arg
                  5
Glu Ser Arg
<210> 211
<211> 15
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: Arbitrary
      peptide
 <400> 211
 Ser Ser Asn His Gln Ser Ser Arg Leu Ile Glu Leu Leu Ser Arg
 <210> 212
 <211> 14
 <212> PRT
 <213> Artificial Sequence
 <220>
 <223> Description of Artificial Sequence: Arbitrary
       peptide
 <400> 212
 Ser Ala Pro Arg Ala Thr Ile Ser His Tyr Leu Met Gly Gly
                    5
   1 .
  <210> 213
  <211> 15
  <212> PRT
  <213> Artificial Sequence
  <220>
  <223> Description of Artificial Sequence: Arbitrary
        peptide
  <400> 213
  Ser Ser Trp Asp Met His Gln Phe Phe Trp Glu Gly Val Ser Arg
                                   10
  <210> 214
  <211> 19
  <212> . PRT
  <213> Artificial Sequence
  <220>
  <223> Description of Artificial Sequence: Arbitrary
```

```
peptide
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Ser Arg Leu Pro Pro Ser Val Phe Ser Met Cys Gly Ser Glu Val Cys <400> 214 5

Leu Ser Arg

<210> 215

<211> 15

<212> PRT

<213> Artificial Sequence

<223> Description of Artificial Sequence: Arbitrary peptide

Ser Ser Pro Gly Ser Arg Glu Trp Phe Lys Asp Met Leu Ser Arg 10 5

<210> 216

<211> 15

<212> PRT

<213> Artificial Sequence

<223> Description of Artificial Sequence: Arbitrary peptide

<400> 216 Ser Ser Glu Tyr Cys Phe Tyr Trp Asp Ser Ala His Cys Ser Arg 15 10 1

<210> 217

<211> 15

<212> PRT

<213> Artificial Sequence

<223> Description of Artificial Sequence: Arbitrary peptide

<400> 217

Ser Ser Leu Thr Ser Arg Asp Phe Gly Ser Trp Tyr Ala Ser Arg 10 5

<210> 218

<211>. 15

<212> PRT

<213> Artificial Sequence

<220>

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<223> Description of Artificial Sequence: Arbitrary
     peptide
Ser Arg Thr Trp Glu Ser Pro Leu Gly Thr Trp Glu Trp Ser Arg
                  5
<210> 219
<211> 15
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Arbitrary
     peptide
Ser Arg Glu Trp Glu Asp Gly Phe Gly Gly Arg Trp Leu Ser Arg
 <210> 220
 <211> 19
 <212> PRT
 <213> Artificial Sequence
 <223> Description of Artificial Sequence: Arbitrary
      peptide
 <400> 220
 Ser Ser Leu Asp Leu Ser Gln Phe Pro Met Thr Ala Ser Phe Leu Arg
 Glu Ser Arg
 <210> 221
 <211> 19
 <212> PRT
 <213> Artificial Sequence
  <223> Description of Artificial Sequence: Arbitrary
       peptide
  Ser Ser Glu Ala Cys Val Gly Arg Trp Met Leu Cys Glu Gln Leu Gly
  <400> 221
   1
  Val Ser Arg
  <210> 222
<211> 15
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<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: Arbitrary
      peptide
Ser Arg Ala Gly Leu Leu Ser Asp Leu Leu Glu Gly Lys Ser Arg
<400> 222
                  5
<210> 223
<211> 15
 <212> PRT
 <213> Artificial Sequence
 <223> Description of Artificial Sequence: Arbitrary
      peptide
 Ser Ser Arg Ser Leu Leu Arg Asp Leu Leu Met Val Asp Ser Arg
                   5
 <210> 224
 <211> 15
 <212> PRT
 <213> Artificial Sequence
 <223> Description of Artificial Sequence: Arbitrary
       peptide
  Ser Ser Asn Lys Leu Leu Tyr Asn Leu Leu Lys Met Glu Ser Arg
  <400> 224
                                       10
   1
  <210> 225
  <211> 15
  <212> PRT
  <213> Artificial Sequence
  <223> Description of Artificial Sequence: Arbitrary
        peptide
  Ser Ser Lys Ser Leu Leu Leu Asn Leu Leu Ser Thr Pro Ser Arg
                                        10
                     5
   <210> 226
   <211> 16
   <212> PRT
   <213> Artificial Sequence
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<220>
<223> Description of Artificial Sequence: Arbitrary
     peptide
His Ser Phe Pro Arg Glu Ser Leu Leu Val Arg Leu Leu Gln Gly Gly
<210> 227
<211> 15
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: Arbitrary
      peptide
Ser Arg Leu Glu Met Leu Leu Arg Ser Glu Thr Asp Phe Ser Arg
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 <210> 228
 <211> 15
 <212> PRT
 <213> Artificial Sequence
 <223> Description of Artificial Sequence: Arbitrary
     peptide
 Ser Arg Leu Glu Glu Leu Leu Lys Trp Gly Ser Val Thr Ser Arg
                                      10
 1
 <210> 229
 <211> 15
 <212> PRT
 <213> Artificial Sequence
 <220>
  <223> Description of Artificial Sequence: Arbitrary
       peptide
  <400> 229
  Ser Arg Leu Glu Gln Leu Leu Lys Glu Glu Phe Ser Tyr Ser Arg
                                        10
                    5
  <210> 230
  <211> .15
  <212> PRT
  <213> Artificial Sequence
  くて20>
```

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<223> Description of Artificial Sequence: Arbitrary
     peptide
<400> 230
Ser Arg Leu Glu Gln Leu Leu Arg Ser Glu Pro Asp Phe Ser Arg
                                     10
                5
<210> 231
<211> 15
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Arbitrary
     peptide
<400> 231
Ser Arg Leu Glu Asp Leu Leu Arg Ala Pro Phe Thr Thr Ser Arg
                                      10
                   5
 <210> 232
 <211> 15
 <212> PRT
 <213> Artificial Sequence
 <223> Description of Artificial Sequence: Arbitrary
      peptide
 <400> 232
 Ser Arg Leu Glu Ser Leu Leu Arg Phe Gly Gln Leu Asp Ser Arg
                                                           15
                                       10
                   5
 <210> 233
 <211> 15
 <212> PRT
 <213> Artificial Sequence
 <220>
 <223> Description of Artificial Sequence: Arbitrary
       peptide
 <400> 233
 Ser Ser Arg Leu Leu Ser Leu Leu Val Gly Asp Phe Asn Ser Arg
                                       10
                   · 5
  <210> 234
  <211> 15
  <212> PRT
  <213> Artificial Sequence
  <220>
  <223> Description of Artificial Sequence: Arbitrary
        Poperde
```

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```
Ser Arg Leu Glu Glu Leu Leu Gly Thr Asn Arg Asp Ser Arg
                 5
<210> 235
<211> 15
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: Arbitrary
     peptide
Ser Arg Leu Glu Glu Leu Leu Met Asp Phe Trp Arg Ser Arg
                                     10
                  5
 <210> 236
 <211> 15
 <212> PRT
 <213> Artificial Sequence
 <223> Description of Artificial Sequence: Arbitrary
     peptide
 Ser Arg Leu Lys Glu Leu Leu Leu Pro Thr Asp Leu Ser Arg
                                      10
              , 5
 <210> 237
 <211> 15
 <212> PRT
 <213> Artificial Sequence
  <223> Description of Artificial Sequence: Arbitrary
       peptide
  Ser Arg Leu Glu Cys Leu Leu Glu Gly Arg Leu Asn Cys Ser Arg
  <400> 237
                                      10
                   5
  <210> 238
  <211> 15
  <212> PRT
  <213> Artificial Sequence
  <220>
  <223> Description of Artificial Sequence: Arbitrary
        peptide
  <400> 238
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```
Ser Ser Lys Leu Tyr Cys Leu Leu Asp Glu Ser Tyr Cys Ser Arg
                                     10
<210> 239
<211> 15
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Arbitrary
      peptide
<400> 239
Ser Arg Leu Ser Cys Leu Leu Met Gly Phe Glu Asp Cys Ser Arg
                   5
<210> 240
 <211> 16
 <212> PRT
 <213> Artificial Sequence
 <223> Description of Artificial Sequence: Arbitrary
       peptide
 <400> 240
 Ser Ser Lys Leu Ile Arg Leu Leu Thr Ser Asp Glu Glu Leu Ser Arg
                                      10
 <210> 241
 <211> 16
 <212> PRT
 <213> Artificial Sequence
 <220>
. <223> Description of Artificial Sequence: Arbitrary
       peptide
 <400> 241
 Ser Ser Arg Leu Met Glu Leu Leu Gln Glu Gly Gln Gly Trp Ser Arg
                                       10
                   5
 <210> 242
 <211> 15
. <212> PRT
 <213> Artificial Sequence
 <223> Description of Artificial Sequence: Arbitrary
      . peptide
  <400> 242
  Ser Ser Asn His Gln Ser Ser Arg Leu Ile Glu Leu Leu Ser Arg
                                       10
```

```
<210> 243
<211> 15
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: Arbitrary
      peptide
Ser Ser Arg Leu Trp Gln Leu Leu Ala Ser Thr Asp Thr Ser Arg
                                      10
                  5
<210> 244
 <211> 15
 <212> PRT
 <213> Artificial Sequence
 <223> Description of Artificial Sequence: Arbitrary
       peptide .
 Ser Ser Lys Leu Trp Gln Leu Leu Ser Ser Pro Ile Asp Ser Arg
 <210> 245
 <211> 15
 <212> PRT
 <213> Artificial Sequence
 <223> Description of Artificial Sequence: Arbitrary
       peptide
  Ser Arg Leu Val Ala Leu Leu Lys Ser Pro Trp Ser Val Ser Arg
  <400> 245
                                       10
  <210> 246
  <211> 15
  <212> PRT
  <213> Artificial Sequence
  <220>
  <223> Description of Artificial Sequence: Arbitrary
        peptide
  <400> 246
  Ser Ser Asn Ser Met Leu Trp Lys Leu Leu Ala Ala Pro Ser Arg
                                        10
                     5
    1
```

```
<210> 247
<211> 15
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Arbitrary
     peptide
<400> 247
Ser Ser Lys Thr Leu Trp Arg Leu Leu Glu Gly Glu Arg Ser Arg
                                      10
<210> 248
<211> 15
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: Arbitrary
      peptide
<400> 248
Ser Arg Ala Gly Pro Val Leu Trp Gly Leu Leu Ser Glu Ser Arg
<210> 249
<211> 15
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: Arbitrary
      peptide
<400> 249
Ser Arg Ser Pro Ile Leu Thr His Leu Leu Ser Leu Gly Ser Arg
                   5
<210> 250
 <211> 15
 <212> PRT '
 <213> Artificial Sequence
<220>
 <223> Description of Artificial Sequence: Arbitrary
     peptide
 <400> 250
 Ser Ser Thr Gly Ile Leu Trp Lys Leu Leu Thr Ala Glu Ser Arg
 <210> 251
 <211> 15
```

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<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Arbitrary
<400> 251
Ser Ser His Gly Ile Leu Trp Arg Leu Leu Ser Glu Gly Ser Arg
                                    10
<210> 252
<211> 11
<212> PRT
<213> Human steroid receptor coactivator la
<400> 252
Lys Leu Val Gln Leu Leu Thr Thr Thr Ala Glu
                  5
  1
<210> 253
<211> 11
<212> PRT
<213> Human steroid receptor coactivator la
              • •
 Ile Leu His Arg Leu Leu Gln Glu Gly Ser Pro
                 5
 <210> 254
 <211> 11
 <212> PRT
 <213> Human steroid receptor coactivator la
 <400> 254
 Leu Leu Arg Tyr Leu Leu Asp Lys Asp Glu Lys
                  5
 <210> 255
 <211> 8.
 <212> PRT
 <213> Human steroid receptor coactivator la
 <400> 255
 Leu Leu Gln Gln Leu Leu Thr Glu
 <210> 256
 <211>,11
  <212> PRT
 <213> Mouse cAMP-responsive element (CREB)-binding protein
  <400> 256
```

```
Gln Leu Ser Glu Leu Leu Arg Gly Gly Ser Gly
                 5
<210> 257
<211> 11
<212> PRT
<213> Mouse cAMP-responsive element (CREB)-binding protein
<400> 257
Gln Leu Val Leu Leu His Ala His Lys Cys
<210> 258
<211> 11
<213> Mouse cAMP-responsive element (CREB)-binding protein
<212> PRT
<400> 258
Tyr Leu Glu Gly Leu Leu Met His Gln Ala Ala
                  5
  1
 <210> 259
 <211> 11
 <212> PRT
 <213> Mouse cAMP-responsive element (CREB)-binding protein
 Leu Leu Ala Ser Leu Leu Gln Ser Glu Ser Ser
                   5
 <210> 260
 <211> 11
 <212> PRT
 <213> Mouse cAMP-responsive element (CREB)-binding protein
 <400> 260
 His Leu Lys Thr Leu Leu Lys Lys Ser Lys Val
                                       10
 <210> 261
  <211> 11
  <212> PRT
  <213> Human RIP140
  <400> 261
  Gln Leu Ala Leu Leu Leu Ser Ser Glu Ala His
                    5
    1
  <210> 262
  <211> 11
  <212> PRT
<Z13> Human RIP140
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```
<400> 262
Leu Leu Leu His Leu Leu Lys Ser Gln Thr Ile
              5
<210> 263
<211> 11
<212> PRT
<213> Human RIP140
<400> 263
Leu Leu Gln Leu Leu Gly His Lys Asn Glu
1 5 10
<210> 264
<211> 11
<212> PRT
<213> Human RIP140
 <400> 264
Val Leu Gln Leu Leu Leu Gly Asn Pro Lys Gly
               5
                                 10
 1
 <210> 265
 <211> 11
 <212> PRT
 <213> Human RIP140
 <400> 265
 Leu Leu Ser Arg Leu Leu Arg Gln Asn Gln Asp
             5
 1
 <210> 266
 <211> 11
 <212> PRT
 <213> Human RIP140
<400> 266
. Val Leu Lys Gln Leu Leu Ser Glu Asn Cys
  1 5
 <210> 267
 <211> 14
 <212> PRT
<213> Artificial Sequence
 <223> Description of Artificial Sequence: Arbitrary
      .peptide
  <400> 267
  Ser Ser Asn His Gln Ser Arg Leu Ile Glu Leu Leu Ser Arg
```

```
<210> 268
<211> 19
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Arbitrary
      peptide
His Val Tyr Gln His Pro Leu Leu Ser Leu Leu Ser Ser Glu His
                                      10
                   5
 Glu Ser Gly
 <210> 269
 <211> 19
 <212> PRT
 <213> Artificial Sequence
 <220>
 <223> Description of Artificial Sequence: Arbitrary
      pepride
 <400> 269
 His Val Glu Met His Pro Leu Leu Met Gly Leu Leu Met Glu Ser Gln
                                     10
                  5
 Trp Gly Ala
 <210> 270
  <211> 19
  <212> PRT
  <213> Artificial Sequence
  <223> Description of Artificial Sequence: Arbitrary
        peptide
  <400> 270
  Gln Glu Ala His Gly Pro Leu Leu Trp Asn Leu Leu Ser Arg Ser Asp
    1
  Thr Asp Trp
  <210>.271
  <211> 19
  <212> PRT
  <213> Artificial Sequence
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<400> 274

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```
<223> Description of Artificial Sequence: Arbitrary
     peptide
Gly His Glu Pro Leu Thr Leu Leu Glu Arg Leu Leu Met Asp Asp Lys
Gln Ala Val
<210> 272
<211> 19
<212> PRT
<213> Artificial Sequence
 <223> Description of Artificial Sequence: Arbitrary
      peptide
 Leu Pro Tyr Glu Gly Ser Leu Leu Leu Lys Leu Leu Arg Ala Pro Val
                                      10
  1
 Glu Glu Val
 <210> 273
 <211> 19
 <212> PRT
 <213> Artificial Sequence
 <223> Description of Artificial Sequence: Arbitrary
       peptide
 <400> 273
 Ser Gly Trp Glu Asn Ser Ile Leu Tyr Ser Leu Leu Ser Asp Arg Val
                              . 10
                   5
  Ser Leu Asp
  <210> 274
  <211> 19
  <212> PRT
  <213> Artificial Sequence
  <223> Description of Artificial Sequence: Arbitrary
       peptide
```

Ala His Gly Glu Ser Ser Leu Leu Ala Trp Leu Leu Ser Gly Glu Tyr

```
Ser Ser Ala
```

```
<210> 275
<211> 19
<212> PRT
<213> Artificial Sequence
```

<220>
<223> Description of Artificial Sequence: Arbitrary peptide

<400> 275
Gly Val Phe Cys Asp Ser Ile Leu Cys Gln Leu Leu Ala His Asp Asn
10
15

Ala Arg Leu

```
<210> 276
<211> 19
<212> PRT
<213> Artificial Sequence
```

<220>
<223> Description of Artificial Sequence:Arbitrary
 peptide

<400> 276
His His Asn Gly His Ser Ile Leu Tyr Gly Leu Leu Ala Gly Ser Asp
1 10 15

Ala Pro Ser

```
<210> 277
<211> 19
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence:Arbitrary
peptide
```

<400> 277
Leu Gly Glu Arg Ala Ser Leu Leu Asp Met Leu Leu Arg Gln Glu Asn
10 15

Pro Ala Trp

<210> 278 <211> 19

```
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: Arbitrary
     peptide
<400> 278
Ser Gly Trp Asn Glu Ser Thr Leu Tyr Arg Leu Leu Gln Ala Asp Ala
                                      10
Phe Asp Val
<210> 279
<211> 19
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Arbitrary
      peptide
 <400> 279
 Pro Ser Gly Gly Ser Ser Val Leu Glu Tyr Leu Leu Thr His Asp Thr
                                      10
 Ser Ile Leu
 <210> 280
 <211> 19
 <212> PRT
 <213> Artificial Sequence
 <220>
 <223> Description of Artificial Sequence: Arbitrary
       peptide
 <400> 280
 Gly Ser Glu Pro Lys Ser Arg Leu Leu Glu Leu Ser Ala Pro Val
                                      10
 Thr Asp Val
 <210> 281
 <211> 19
 <212> PRT
  <213> Artificial Sequence
  <223> Description of Artificial Sequence: Arbitrary
       peptide
```

```
<400> 281
His Pro Thr His Ser Ser Arg Leu Trp Glu Leu Leu Met Glu Ala Thr
15
1
```

Pro Thr Met

<210> 282 <211> 19 <212> PRT <213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Arbitrary peptide

Leu Leu Thr

<210> 283 <211> 19 <212> PRT <213> Arrificial Sequence

<400> 283
Trp Glu Glu His Ser Gln Met Leu Leu His Leu Leu Asp Thr Gly Glu
15

Ala Val Trp

<210> 284 <211> 19. <212> PRT <213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:Arbitrary
 peptide

<400> 284
Pro Val Gly Glu Pro Gly Leu Leu Trp Arg Leu Leu Ser Ala Pro Val
1 10 15

Glu Arg Glu

,F

```
<210> 285
<211> 12
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: Arbitrary
      peptide
Met Ser Trp Tyr Glu Phe Met Thr Glu Glu Ser Met
<400> 285
                  5
 <210> 286
 <211> 15
 <212> PRT
 <213> Artificial Sequence
 <223> Description of Artificial Sequence: Arbitrary
 <220>
      peptide
 Ala Lys His Asp Leu Ser Trp Tyr Glu Phe Leu Gln Leu Pro Ile
                                      10
 <210> 287
 <211> 17
 <212> PRT
 <213> Artificial Sequence
  <220>
  <223> Description of Artificial Sequence: Arbitrary
       peptide
  Ser Arg Leu Ser Trp Trp Glu Phe Leu Gly Ala Ser Asp Cys Gly Thr
 <400> 287
                                 10 15
                   5
  Cys
  <210> 288
  <211> 11
  <212> PRT
  <213> Artificial Sequence
  <223> Description of Artificial Sequence: Arbitrary
       peptide
   <400> 288
  Asp Leu Leu Ser Leu Lys Glu Phe Leu Ala Thr
                    5
```

```
<210> 289
<211> 13
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: Arbitrary
    peptide
<400> 289
Ser Ser Pro Asn Leu Leu Thr Leu Glu Glu Phe Leu Ser
                 5
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IN THE UNITED STATES PATENT AND TRADEMARK OFFICE

In re Application of:) ART UNIT: 1627
PAIGE et al	Examiner: T. Wessendorf
Appln. No.: 09/429,331) Washington, D.C.
Filed: October 28, 1999) February 27, 2001
For: METHOD OF PREDICTING THE ABILITY OF COMPOUNDS TO MODULATE THE BIOLOGICAL ACTIVITY OF RECEPTORS	Atty.Docket: PAIGE=1D)))

SUPPLEMENTAL RESPONSE TO "SEQUENCE LISTING" REQUIREMENT

Honorable Commissioner of Patents Washington, D.C. 20231

Sir:

Supplementing the response filed February 20, 2001, please further amend the application as follows:

IN THE SPECIFICATION

At page 162, line 34, replace "K = C or T" with --K = G or T--.

Please replace the present page 237 with the enclosed substitute page, which provides SEQ ID NOs:316-360 for the sequences on this page. If there was no page numbered "237" in the application as filed, please insert this page between pages "236" and "238".

The enclosed "Sequence Listing" pages 1, 4 and 79-90 replace pages 1, 4 and 79 of the "Sequence Listing" submitted February 20, 2001.

In re Appln. No-09/429,331

REMARKS

1. At the time the February 20, 2001, response was prepared, counsel's file copy of the specification was missing page 237. Hence, 'the sequences appearing on that page were not incorporated into the Sequence Listing filed on that date.

Since counsel received a postcard receipt (copy enclosed) acknowledging the filing of a 293 page specification, counsel assumes that page 237 was missing only from counsel's file copy and not from the original filed with the PTO.

If counsel is mistaken, inserting page 237 at this time does not constitute the addition of "new matter". At page 1, lines 3-10, it is stated:

This application is a continuation-in-part of PCT/US99/06664, filed March 26, 1999, which is a continuation-in-part of 60/115,345, filed January 8, 1999, which is a continuation-in-part of Paige et al., Serial No. 60/099,656, filed September 9, 1998, which is a continuation-in-part of Paige et al., Serial No. 60/082,756, filed April 23, 1998. All of the above applications are hereby incorporated-by-reference.

Page 237 of this application sets forth Table 1, and part of Table 2. It is identical to page 152 of the above-identified, incorporated-by-reference PCT application. Hence, even if inadvertently omitted from this application as filed, it can be provided without adding "new matter".

In re Appln. No 39/429,331

- 2. At page 162, we correct an obvious typographical error in the identification of ambiguous nucleotide "K", which denotes "G" or "T", not "C" or "T". See MPEP \$2422, page 2400-20, Table 1. The NNK codon, specified at page 162, line 33, encodes all 20 amino acids. If the third position were C/T (Y), instead of G/T (K), then Met (ATG), Trp (TGG), Ser (TCA, TCG), Gln (CAA, CAG), Lys (AAA, AAG) and Gly (GAA, GAG) would not be encoded, inconsistent with the identification of X in LXXLL (page 162, line 29) as "any AA". This error was also corrected on page 4 of the Sequence Listing at <223> in SEQ ID NO:14.
 - 3. Applicants hereby submit the following:
 - [XX] an amendment to the paper copy of the "Sequence Listing" submitted on February 20, 2001, the amendment being in the form of substitute pages 1 and 79 and new pages 80-90;
 - [XX] the Sequence Listing in computer readable form, complying with \$1.821(e) and \$1.824, including, if an amendment to the paper copy is submitted, all previously submitted data with the amendment incorporated therein;
- [XX] 4. The description has been amended to comply with \$1.821(d).

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5. The undersigned attorney or agent hereby states as follows:

- (a) this submission is not believed to include new matter [\$1.821(g)];
- (b) the contents of the paper copy (as amended, if applicable) and the computer readable form of the Sequence Listing, are believed to be the same [\$1.821(f) and \$1.825(b)];
- (c) if the paper copy has been amended, the amendment is believed to be supported by the specification and is not believed to include new matter [\$1.825(a)]; and

Respectfully submitted,

BROWDY AND NEIMARK

Attorneys for Applicant(s)

By:

Iver P. Cooper

Registration No. 28,005

IPC:al
624 Ninth Street, N.W.
Washington, D.C. 20001
Telephone No.: (202) 628-5197
Facsimile No.: (202) 737-3528
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Enclosures:

Paper Sequence Listing pp. 1,4 and 79-90 Substitute CRF Substitute page 237 Page 152 of PCT/US99/06664 Copy of stamped postcard receipts

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PCT/US99/06664 '

Table 1

Peptides the Bind to the Unliganded (unactivated)
Estrogen Receptor

	Eschogen	, Macop	Sequence	Phage #
5	S R S A	WESPLG	TWEWSR HYLMGG FPWGVSR	4 48 1
·	S S S S S S	W D R L S D W I R L R D	FPWGVSR LPWGESR	2 3 31
10	. s s	WVVLRD	L P W G S R L P W G S R C S G L W S R	29 · 7
	SS	B G I C F F W R N L C F F W	DDEYCSR	35 41 47
15.	H H	ннкнран	PHTYGG	

Table 2

Peptides that Bind to the Estradiol Activated

	Receptor	•
	Sequence	Phage #
		1/2
20	SKAGILLI	б
	5 5 K 5 2 2 K 1 6 B	22
		23
	22777777	42
		3
25	5 6 7 7 7 7 7 7 7 7 7 7 7 7 7 7 7 7 7 7	11
		21
	2 1 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2	. 27
•		28
	3 1 1 1 2 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1	29
30		19/20
		30
	2 4 4 4 4 4 4 4 4 4 4 4 4 4 4 4 4 4 4 4	15
		34
		35
35	224	36
	5 K H 5 C 2 E 4	37
	SSKLIRLLTSDEELSR	40
	SSRUMBULZIO	4
	SSNHQSSRLIELLSR	16
40	SSRLWQLLASTDTSR	13/14
	SSNSMLWKLLAAPSR	17
-	SSKTLWRLLEGERSR	32
	SRAGPVLWGLLSESR	5.2 . 5
	SSLTSRDFGSWYASR	
45	SSWVRLSDFPWGVSR	24/25
• •	SSEYCFYDSAHCSR	33 .
	SRSLLECHLMGNCSR	7
	SSELLRWHLTRDTSR	8
	S R L E Y W L K W E P G P S R	. 12
E 0	SRSDSILWRMLSESR	31
50	SSKGVLWRMLAEPVSR	38/39
	HSHGPLTLNLLRSSGG	41
	SSAGGGAPAGSTPSR	26

237

Table 1

From-BROWDY NEIMARK

Peptides	the	Bind	to	the	Unliganded	(unactivat	(ba:
Estrogen Recep							5 2

		_	Sequence	sea id no:	Phage #
_	c n	WESPLG	TWEWSR	316	4
5	s a		HYLMGG	317	48
	8 8 8 8		FPWGVSR	318	1
	5 S	N	FPWGVSR	319	2
	5 S	, , , , , , , , , , , , , , , , , , , ,	LPWGESR	320	3
	S S	M # 12 = 20 = 2		321	31
10	\$ 5 S 5		~ ~ ~ ~ ~ ~	322	29
•	-			323	7
				324	. 35
		SNLCFFW		325	41
			PHTYGG	326	47
1.5	. н		-		

Table 2

Peptides that Bind to the Estradiol Activated Receptor

	~	SEB IDNO:	Phage #
	Sequence		
20	SRAGLLSDLLEGKSR	327	1/2
20		328	6
	CCNKILYNLLKMESR	329	22
	SSKSLLLNLLSTPSR	330	23
	HSFPRESLLVRLLQGG	331	42
		332	3
25	S A L L L L L L L L L L L L L L L L L L	333	11
	SKALLERBBOVED	334	21
	SKIIIX	335	. 27
		336	28
	SKHADIA	337	29
30		338	19/20
	S S C I I S S C B C B		30
	SKUMBALL	339 340	15
	SKIKIIII	341	· 34
	SRUBCHARO	345	35
35	55441044	343	36
	2 K H D C D T T C B	3 (3 3 (4	37
	552442444	3 4 5	40
	SSRLMELLQEGQGWSR	34 P	4
	SSNHQSSRLIELLSR		16
40	SSRLWQLLASTDTSR	347	13/14
••	SSNSMLWKLLAAPSK	348	17
	SSKTLWRLLEGERSR	349	32
	SRAGPVLWGLLSESR	350	5
	SSLTSRDFGSWYASR	351	24/25
45	SSWVRLSDFPWGVSR	352	
40	SSEYCFYDSAHCSR	353	33
	GRSLLECHLMGNCSR	3 <i>5</i> ¥	7
	SSELLRWHLTRDTSR	3 <i>5</i> 5	8
	SRLEYWLKWEPGPSR	356	12
		357	31
50	S R S D S T T T T T T T T T C D	358	38/39
	552444444	359	41
	n and 1 2 1 1 1 1 1 C B	3 Lo	26
	SSAGGGAPAGSTPSR	-	

SEQUENCE LISTING

- <110> PAIGE, Lisa A. MCDONNELL, Donald P. CHANG, Ching Yu NORRIS, John HAMILTON, Paul T. FOWLKES, Dana M. BARNETT, Tom CHRISTIANSEN, Dale J. BUEHRER, Benjamin
- <120> METHOD OF PREDICTING THE ABILITY OF COMPOUNDS TO MODULATE THE BIOLOGICAL ACTIVITY OF RECEPTORS
- <130> PAIGEID
- <140> 09/429,331
- <141> 1999-10-28
- <150> PCT/US99/06664
- <151> 1999-03-26
- <150> 60/082,756
- <151> 1998-04-23
- <150> 60/099,656
- <151> 1998-09-09
- <150> 60/115,345
- <151> 1999-01-08
- <160> 360
- <170> PatentIn Ver. 2.0
- <210> 1
- <211> 20
- <212> PRT
- <213> Arcificial Sequence
- <223> Description of Artificial Sequence: Arbitrary peptide
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- Gly Ser Gly Lys
- <210> 2
- <211> 8
- <212> PRT
- <213> Artificial Sequence

<400> 11

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Ser Ser Trp Asp Met His Gln Phe Phe Trp Glu Gly Val Ser Arg
<210> 12
<211> 15
<212> PRT
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<223> Description of Artificial Sequence: Arbitrary
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Ser Ser Pro Gly Ser Arg Glu Trp Phe Lys Asp Met Leu Ser Arg
                                      10
<210> 13
<211> 15
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<213> Artificial Sequence
<223> Description of Artificial Sequence: Arbitrary
     peptide
<400> 13
Ser Ser Leu Thr Ser Arg Asp Phe Gly Ser Trp Tyr Ala Ser Arg
<210> 14
<211> 88
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<223> Description of Artificial Sequence: DNA encoding random
      peptide library of Ex. 101.1
<220>
<223> N at each occurrence is A, C, G or T; K at each
      occurrence is G or T
<400> 14
agtgtgtgcc tegagannkn nknnknnknn knnknnketg nnknnketge tgnnknnknn 60
knnknnknnk nnktetagae tgtgeagt
<210> 15
<211> 15
<212> DNA
<213> Artificial Sequence
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<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: Arbitrary
      peptide
<400> 313
Ser Ser His Trp Ser Ser Asp Ser Ile Phe Pro Gly Phe Trp Tyr Ser
                                      10
Gly
<210> 314
<211> 13
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Arbitrary
     peptide
<400> 314
Ser Arg Gly Gly Val Asp Leu Asp Ile Gly Asn Ser Ala
<210> 315
<211> 11
 <212> PRT
 <213> Artificial Sequence
 <220>
 <223> Description of Artificial Sequence: Arbitrary
       pepride
 <400> 315
 Glu Gly Glu Asp Val Arg Thr Arg Ile Ala Asn
   1
 <210> 316
 <211> 14
 <212> PRT
 <213> Artificial Sequence
 <223> Description of Artificial Sequence: Arbitrary
       peptide
 <400> 316
 Ser Arg Trp Glu Ser Pro Leu Gly Thr Trp Glu Trp Ser Arg
                                       10
                   5
 <210> 317
 <211> 14
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<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Arbitrary
      peptide
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<223> Description of Artificial Sequence: Arbitrary
<400> 318
Ser Ser Trp Val Arg Leu Ser Asp Phe Pro Trp Gly Val Ser Arg
                                                           15
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<211> 15
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Arbitrary
      peptide
<400> 319
Ser Ser Trp Asp Arg Leu Ser Asp Phe Pro Trp Gly Val Ser Arg
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 <223> Description of Artificial Sequence: Arbitrary
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 Ser Ser Trp Ile Arg Leu Arg Asp Leu Pro Trp Gly Glu Ser Arg
                   5
 <210> 321
 <211> 14
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 <213> Artificial Sequence
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<220>

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<223> Description of Artificial Sequence: Arbitrary
     peptide
<400> 321
Ser Ser Trp Val Leu Leu Arg Asp Leu Pro Trp Gly Ser Arg
                                      10
                  5
<210> 322 ·
<211> 14
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<223> Description of Artificial Sequence: Arbitrary
   peptide
<400> 322
Ser Ser Trp Val Val Leu Arg Asp Leu Pro Trp Gly Ser Arg
                  5
<210> 323
<211> 15
<212> PRT
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      peptide
<400> 323
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 <210> 324
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 <400> 324
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<223> Description of Artificial Sequence: Arbitrary
     peptide
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                                     10
<210> 326
<211> 14
<212> PRT
<213> Artificial Sequence
<220>
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    · peptide
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<210> 327
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<400> 327
Ser Arg Ala Gly Leu Leu Ser Asp Leu Leu Glu Gly Lys Ser Arg
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                                      10
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      peptide
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                                       10
 <210> 329
 <211> 15
 <212> PRT
 <213> Artificial Sequence
```

<223> Description of Artificial Sequence: Arbitrary

peptida

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<400> 329
 Ser Ser Asn Lys Leu Leu Tyr Asn Leu Leu Lys Met Glu Ser Arg
                                       10
   1 ·
 <210> 330
 <211> 15
 <212> PRT
 <213> Artificial Sequence
 <223> Description of Artificial Sequence: Arbitrary
      peptide
· <400> 330
 Ser Ser Lys Ser Leu Leu Leu Asn Leu Leu Ser Thr Pro Ser Arg
                                       10
 <210> 331
 <211> 16
 <212> PRT
 <213> Artificial Sequence
 <220>
 <223> Description of Artificial Sequence: Arbitrary
       peptide
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 His Ser Phe Pro Arg Glu Ser Leu Leu Val Arg Leu Leu Gln Gly Gly
                  5
 <210> 332
 <211> 15
 <212> PRT
 <213> Artificial Sequence
 <223> Description of Artificial Sequence: Arbitrary
       peptide
 <400> 332
 Ser Arg Leu Glu Met Leu Leu Arg Ser Glu Thr Asp Phe Ser Arg
                                       10
                    5
 <210> 333
 <211> 15
  <212> PRT
 <213> Artificial Sequence
 <220>
  <223> Description of Artificial Sequence: Arbitrary
       peptide
  <400> 333
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. Ser Arg Leu Glu Glu Leu Leu Lys Trp Gly Ser Val . Thr Ser Arg
                                       10
 <210> 334
 <211> 15
 <212> PRT
 <213> Artificial Sequence
 <220>
 <223> Description of Artificial Sequence: Arbitrary
       peptide
 <400> 334
 Ser Arg Leu Glu Gln Leu Leu Lys Glu Glu Phe Ser Tyr Ser Arg
                                       10
 <210> 335
 <211> 15
 <212> PRT
 <213> Artificial Sequence
 <220>
 <223> Description of Artificial Sequence: Arbitrary
       peptide
 <400> 335
 Ser Arg Leu Glu Gln Leu Leu Arg Ser Glu Pro Asp Phe Ser Arg
                   <sub>.</sub>5
                                        10
   1
 <210> 336
 <211> 15
 <212> PRT
 <213> Artificial Sequence
 <223> Description of Artificial Sequence: Arbitrary
       peptide
 <400> 336
 Ser Arg Leu Glu Asp Leu Leu Arg Ala Pro Phe Thi Thr Ser Arg
   1
 <210> 337
 <211> 15
 <212> PRT
 <213> Artificial Sequence
 <220>
 <223> Description of Artificial Sequence: Arbitrary
       peptide
 <400> 337.
  Ser Arg Leu Glu Ser Leu Leu Arg Phe Gly Gln Leu Asp Ser Arg
                    5
                                        10
                                                             15
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<210> 338
<211> 15
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: Arbitrary
     peptide
<400> 338
Ser Ser Arg Leu Leu Ser Leu Leu Val Gly Asp Phe Asn Ser Arg
                                     10
                 5
<210> 339
<211> 15
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Arbitrary
      peptide
<400> 339
Ser Arg Leu Glu Glu Leu Leu Gly Thr Asn Arg Asp Ser Arg
                  5
                                      10
<210> 340
<211> 15
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Arbitrary
      peptide
<400> 340°
Ser Arg Leu Lys Glu Leu Leu Leu Pro Thr Asp Leu Ser Arg
                , 5
<210> 341
 <211> 15
 <212> PRT
 <213> Artificial Sequence
 <223> Description of Artificial Sequence: Arbitrary
      peptide
 <400> 341
 Ser Arg Leu Glu Cys Leu Leu Glu Gly Arg Leu Asn Cys Ser Arg
```

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<210> 342
<211> 15
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<223> Description of Artificial Sequence: Arbitrary
      peptide
<400> 342
Ser Ser Lys Leu Tyr Cys Leu Leu Asp Glu Ser Tyr Cys Ser Arg
                                      10
<210> 343
<211> 15
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Arbitrary
      peptide
<400> 343
Ser Arg Leu Ser Cys Leu Leu Met Gly Phe Glu Asp Cys Ser Arg
<210> 344
<211> 16
<212> PRT
<213> Artificial Sequence
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      peptide
Ser Ser Lys Leu Ile Arg Leu Leu Thr Ser Asp Glu Glu Leu Ser Arg
                  5
                                     10
<210> 345
<211> 16
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Arbitrary
      peptide
<400> 345
Ser Ser Arg Leu Met Glu Leu Leu Gln Glu Gly Gln Gly Trp Ser Arg
                                      10
<210> 346
<211> 15
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<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: Arbitrary
     peptide
<400> 346
Ser Ser Asn His Gln Ser Ser Arg Leu Ile Glu Leu Leu Ser Arg
                                      10
<210> 347
<211> 15
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<213> Artificial Sequence
<223> Description of Artificial Sequence: Arbitrary
      peptide
<400>,347
Ser Ser Arg Leu Trp Gln Leu Leu Ala Ser Thr Asp Thr Ser Arg
<210> 348
<211> 15
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Arbitrary
      peptide
<400> 348
Ser Ser Asn Ser Met Leu Trp Lys Leu Leu Ala Ala Pro Ser Arg
                  5
                                      10
<210> 349
<211> 15
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: Arbitrary
      peptide
<400> 349
Ser Ser Lys Thr Leu Trp Arg Leu Leu Glu Gly Glu Arg Ser Arg
                                      10
<210> 350
<211> 15
<212> PRT
<213> Artificial Sequence
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<223> Description of Artificial Sequence: Arbitrary
      peptide
<400> 350
Ser Arg Ala Gly Pro Val Leu Trp Gly Leu Leu Ser Glu Ser Arg
<210> 351
<211> 15
<212> PRT
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<223> Description of Artificial Sequence: Arbitrary
      peptide
<400> 351
ser Ser Leu Thr Ser Arg Asp Phe Gly Ser Trp Tyr Ala Ser Arg
                                      10
<210> 352
<211> 15
<212> PRT
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<220>
<223> Description of Artificial Sequence: Arbitrary
      peptide
<400> 352
Ser Ser Trp Val Arg Leu Ser Asp Phe Pro Trp Gly Val Ser Arg
                  5
<210> 353
<211> 14
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Arbitrary
      peptide
<400> 353
Ser Ser Glu Tyr Cys Phe Tyr Asp Ser Ala His Cys Ser Arg
<210> 354
<211> 15
<212> PRT
<213> Artificial Sequence
<220>
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<223> Description of Artificial Sequence: Arbitrary
     peptide
<400> 354
Ser Arg Ser Leu Leu Glu Cys His Leu Met Gly Asn Cys Ser Arg
<210> 355
<211> 15
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: Arbitrary
    . peptide
<400> 355
Ser Ser Glu Leu Leu Arg Trp His Leu Thr Arg Asp Thr Ser Arg
<210> 356
<211> 15
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Arbitrary
      peptide
<400> 356
Ser Arg Leu Glu Tyr Trp Leu Lys Trp Glu Pro Gly Pro Ser Arg
  1
<210> 357
<211> 15
<212> PRT
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<223> Description of Artificial Sequence: Arbitrary
      peptide
<400> 357
Ser Arg Ser Asp Ser Ile Leu Trp Arg Met Leu Ser Glu Ser Arg
                                      10
<210> 358
<211> 16
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Arbitrary
      peptide
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<400> 358
Ser Ser Lys Gly Val Leu Trp Arg Met Leu Ala Glu Pro Val Ser Arg
                                     10
<210> 359
<211> 16
<212> PRT
<213> Artificial Sequence -
<220>
<223> Description of Artificial Sequence: Arbitrary
<400> 359
His Ser His Gly Pro Leu Thr Leu Asn Leu Leu Arg Ser Ser Gly Gly
                 5
<210> 360
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     peptide
<400> 360
Ser Ser Ala Gly Gly Gly Ala Pro Ala Gly Ser Thr Pro Ser Arg
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